STIC-Biotech/ChemLib

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Schultz, James

Thursday, May 15, 2003 7:52 AM STIC-Biotech/ChemLib

To: Subject: Seq Search 09/780,929

RECEIVED

MAY 15 2003

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Hello,

Could you please run a length limited nucleotide sequence search on SEQ ID NOS: 97 (15 nt long) and 98 (18 nt long) in the above entitled case, where the maximum size of the returned hit is no longer than 20 nucleotides long? Thanks,

Doug Schultz

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From: Sent:

Schultz, James

Thursday, May 15, 2003 7:53 AM STIC-Biotech/ChemLib

To: Subject:

But wait there's more--Seq Search 09/780,929

MAY 15 2900

The sequence search request below was just submitted--I forgot to include that I also need the interference databases searched. Thanks, Doug Schultz

----Original Message----

From: Sent:

Schultz, James

To:

Thursday, May 15, 2003 7:52 AM

Subject:

STIC-Biotech/ChemLib Seq Search 09/780,929

Hello,

Could you please run a length limited nucleotide sequence search on SEQ ID NOS: 97 (15 nt long) and 98 (18 nt long) in the above entitled case, where the maximum size of the returned hit is no longer than 20 nucleotides long? Thanks,

Doug Schultz

	TYPE OF SEARCH:	VENDOR/COST (where applic.)
Searcher:	NA Sequences:	STN:
Phone:	AA Sequences:	DIALOG:
Location:	Structures:	Questel/Orbit:
Date Picked Up: 5//6	Bibliographic:	DRLink:
Date Completed: 15/25	Litigation:	Lexis/Nexis:
Searcher Prep/Review:	Full text:	Sequence Sys.:
Clerical:	Patent Family:	WWW/Internet:
Online time:	Other:	Other (specify):

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Database :
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Maximum Match 1008
Listing first 45 summaries
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ALIGNMENTS

TITLE JOURNAL	AUTHORS	REFERENCE	ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	LOCUS	AX214295	RESULT 1
Nucleozymes with endonuclease activity Patent: WO 0159102-A 108 16-AUG-2001; RIBOZYME PHARMACEUTICALS, INC. (US); Yale University (US)	Breaker, R. and Emilsson, G.	artificial sequences. 1 (bases 1 to 15)	synthetic construct .	synthetic construct.		AX214295.1 GI:15524372	AX214295	02.	AX214295 15 bp mRNA linear PAT 06-SEP-2001		

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RESULT 2
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Spermatophyta; Magnoliophyta; eudicotyledons; c
Caryophyllidae; Caryophyllales; Chenopodiaceae;
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Caryophylidae; Caryophyllales; Chenopodiaceae; Beta.
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Targeted chromosomal genomic alterations in p
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AX254729
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20hlhhoefer,D., Baeuerle,P., Klein,C. and Neumann,F.:
Identification of modulators of the interferon gamma
pathway and their use in restenosis treatment
Patent: WO 0170953-A 18 27-SEP-2001;
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Mrna amplification
Patent: WO 0171027-A 18 27-SEP-2001;
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UNIVERSITE CATHOLIQUE DE LOUVAIN (BE
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/db_xref="taxon:32630"
/note="oligonucleotide"
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Sequence 40 from Patent W00162911.
AX272471
AX272471.1 GI:16545208
                                                                               Sequence 41 from Patent Ax272472
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Method for qualitative and/or quantitative interactions on probe arrays
Patent: WO 0202810-A 21 10-JAN-2002;
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Eukaryota; Metazoa;
Mammalia; Eutheria;
                          Homo sapiens
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/db_xref="taxon:33630"
/note="0ligonukleotidsonde"
5 c 2 g 6 t
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/db_xref="taxon:9606"
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                                                                    GI:16545209
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Chordata;
Primates;
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Pred. No. 4.6e+05;
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Pred. No. 4.6e+05;
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              Craniata; Vertebrata; Euteleostomi;
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Sequence 222 from Patent W00162911.
AX272653
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AX272654
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             Method and reagent for the inhibition of grid Patent: WO 0162911-A 223 30-AUG-2001; RIBOZYME PHARMACEUTICALS, INC. (US); GLAXO G
                                                  Jarvis,T., von Carlowitz,I., Mcswiggen,J.A., Ellis,J.H.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/db_xref="taxon:9606"
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66.7%;
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Pred. No. 4.6e+05;
3; Mismatches 1;
                                                                                                                                                                                                                                                                                                                  Score 10.4; DB 6; Pred. No. 4.6e+05;
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AX272993/c
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AX299883/c
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AX299883
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                      Conner,T.W., Dubois,P., Malven,M. and Masucci,J.I.
Plant regulatory sequences for selective control
Patent: WO 0183790-A 24 08-NOV-2001;
Monsanto Technology LLC (US)
Location/Qualifiers
                                                                                                                                                                                                                      synthetic construct synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Method and reagent for the inhibition of grid patent: WO 0162911-A 562 30-AUG-2001;
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/db_xref="taxon:9606"
4 c 3 g
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/db_xref="taxon:9606"
5 c 3 g
                                                     /organism="synthetic construct"
/db_xxef="taxon:32630"
/note="Fully Synthesized Primer"
1 6 c 4 g 6 t
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83.3%;
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Pred. No. 4.6e+05;
3; Mismatches 1;
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AUTHORS
TITLE
Search completed: May 22, 2003, 19:36:39 Job time: 876.364 secs
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AX325561
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                                                                                                                       Query Match 69.3%; Score 10.4; DB 6; Length 17; Best Local Similarity 75.0%; Pred. No. 4.6e+05; Matches 9; Conservative 2; Mismatches 1; Indels
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16 GACAACGTGAAG 5
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Targeted Chromosomal genomic alterations in plants using modified single stranded oligonucleotides
Patent: WO 0192512-A 1699 06-DEC-2001;
UNIVERSITY OF DELAWARE (US)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                Beta vulgaris.
Beta vulgaris.
Beta vulgaris.
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Caryophyllidae; Caryophyllales; Chenopodiaceae; Beta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AX325561 17 bp DNA Sequence 1699 from Patent WO0192512. AX325561
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                                                                                                                                                                                                     /organism="Beta vulgaris"
/db_xref="taxon:161934"
0 c 5 g 6
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GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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11.4	11.4	11.4	11.4	11.4	12	12	12	15	Score
76.0	76.0	76.0	76.0	76.0	80.0	80.0	80.0	100.0	Query Match Length DB
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AAZ03321	ABF76055	ABF76054	ABF31425	ABF31424	ABK26344	ABK26343	AAA18478	AAS12347	ID
PCR primer used to	Oligonucleotide SE	Oligonucleotide SE	Oligonucleotide SE	Oligonucleotide SE	Increased starch p	Increased starch p	Human TIE-2 substr	DNA encoding deoxy	Description

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H974 C196 C196	X819 X819 Z698 X323	AAS965 ABL565 AAA801 AAZ628	F061 K263	F017 F017 K985	ABI3269 ABI3491 ABI4530 ABC8990 ABC8990	V036 V036 V939 V923 V923 V923	10 m C C m III
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ALIGNMENTS

RESULT 1 AAS12347

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08-FEB-2000;
31-MAR-2000;
New nucleic acids with endonuclease activity, such as ribozymes and
                                            WPI; 2001-536526/59.
                                                                                 Breaker R,
                                                                                                                                                                                                                                                 08-FEB-2001; 2001WO-US04223
                                                                                                                                                                                                                                                                                                                                   WO200159102-A2.
                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                Deoxyribozyme; cytostatic; endonuclease; RNA cleavage; DNA cleavage;
gene therapy; plant; fungus; bacteria; mammal; ribozyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-NOV-2001 (first entry)
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                                                                                                                         (RIBO-) RIBOZYME PHARM INC.
(UYYA ) UNIV YALE.
                                                                                                                                                                                                                                                                                            16-AUG-2001.
                                                                                 Beigelman L,
                                                                                                                                                                                     2000US-0181360
2000US-0193646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         deoxyribozyme #7.
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                    Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     fungal cell. They are used to cleave a separate nucleic acid, preferably RNA. The nucleic acids are used to inhibit gene expression and/or cell proliferation, and can be used to treat a disease or condition. More than one nucleic acid can be independently targeted to the same or different sites in a cell. The nucleic acids may be used to study DNA. The modifications to the nucleic acids optimises their catalytic activity and can maintain or enhance their activity. They exhibit a high degree of specificity for RNA. The present sequence represents the coding sequence of deoxyribozyme #7 used in the method of the invention.
                                            Claim
                                                                        Novel ribozymes stability of an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human TIE-2 substrate sequence SEQ ID NO:1704.
                                                                                                                                                                                                                       27-MAR-1998;
                                                                                                                                                                                                                                                    24-MAR-1999;
                                                                                                                                                                                                                                                                                                                   WO9950403-A2
                                                                                                                                                                                                                                                                                                                                                                                Kippel-Trenaunay-Weber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA18478
                                                                                                                        WPI; 1999-591315/50
                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA18478 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to nucleic acid molecules with endonuclease activity, which are particularly useful for cleavage of RNA or DNA. The nucleic acids are used in a pharmaceutical composition and are used to modulate expression of a gene in a plant, mammalian, bacterial or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleozymes, for modulat:
bacterial or fungal cell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGAUAACGUGAAGAU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
15; Conser
                                           Page 97;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 77; 96pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 7 A; 1 C; 4 G; 3 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for modulating gene expression in a plant, mammalian, fundal cell .
                                                                                                                                                                                                                     98US-0079678
                                                                                                                                                                                                                                                   99WO-US06507
                                                                  for modulating the synthesis, expression and/or {\tt mRNA} encoding an angiogenic factors -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
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                                                                                                                                                      Jarvis
                                         English
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                          Osler-Weber-Rendu syndrome; ss
                                                                                                                                                    Coeshott C,
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                                                                                                                                                   McSwiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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RESULT 3
ABK26343
ID ABK2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 C gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to CC AAA17167 and AAA17561 to AAA17622 represent tribozyme sequences for ARNT, CC and AAA17168 to AAA17560 and AAA17633 to AAA17684 represent their CC corresponding target sequences; AAA17685 to AAA18385 and AAA19087 to CC AAA19154 represent ribozyme sequences for Tie-2, and AAA18386 to AAA19086 condata and AAA19157 to AAA19157 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      integrin subunit alpha-6, or integrin subunit beta-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 AGAUAACGUGAA 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 20;
. 3.2e+03;
ches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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porphyric herbicide resistance; triazine resistance; disease resistance; modified oil production; modified starch production; waxy starch; altered floral morphology; male-sterile plant; albino mutant; modified fatty acid content; reduced palmitate production; albino plant; increased stearate production; reduced linolenic acid production; Chromosomal genomic alteration; genome altering oligonucleotide; PCR; o-methyl modification; LNA modification; phosphorothicate linkage; 09-APR-2002 photosynthetic amino acid over production; herbicide resistance; glyphosate resistance; imidazolinone herbicide resistance; sulphonylurea herbicide resistance; DNA repair; DNA alteration; environmental tolerance; hygromycin B, abiotic stress tolerance; improved nutritional value; hygromycin; Increased starch production genome altering oligonucleotide #195 ABK26343; (first entry)

primer

ss;

The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl

01-JUN-2000; 30-OCT-2000; 27-MAR-2001;

; 2000US-208538P. ; 2000US-244989P. ; 2001US-0818875.

01-JUN-2001; 06-DEC-2001

2001WO-US17672

Synthetic.

Beta vulgaris

process.

WO200192512-A2

(UYDE) UNIV DELAWARE

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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    more of these modifications. The oligonucleotides are useful for directing repair or alteration of plant genetic information. The oligonucleotides are particularly useful for creating plants with desired phenotypes, e.g. environmental or abiotic stress tolerance, improved nutritional value (e.g. altering amino acid content of plants or conferring amino acid over production), herbicide resistance (e.g. glyphosate resistance, imidazolinone and sulphonylurea herbicide resistance, porphyric herbicide resistance or triazine resistance),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                improved modified
             WO200192512-A2
                                                                                                              altered floral morphology; male-sterile plant; albino mutant; modified fatty acid content; reduced palmitate production; albino increased stearate production; reduced linolenic acid production;
                                                                                                                                                                        modified oil production; modified starch production; waxy starch;
                                                                                                                                                                                                             amino acid over production; herbicide resistance; glyphosate resistance; imidazolihone herbicide resistance; sulphonylurea herbicide resistance;
                                                                                                                                                                                                                                                                      o-methyl modification; LNA modification; phosphorothioate linkage; DNA repair; DNA alteration; environmental tolerance; hygromycin-B;
                                                                                                                                                                                                                                                                                                                  Chromosomal genomic alteration;
                                                                                                                                                                                                                                                                                                                                                  Increased starch production genome altering oligonucleotide #196
                                                                                                                                                                                                                                                                                                                                                                                                 09-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                     ABK26344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disease resistance, modified oil production, modified starch production (e.g. increased starch or production of waxy starch), altered floral morphology (e.g. male-sterile plants) or modified fatty acid content (e.g. reduced palmitate, increased stearate or reduced inolenic acid). The oligonucleotides are also useful for producing albino mutants for the analysis of photosynthetic processes. This sequence represents a genome
                                                                 Beta vulgaris
                                                                                                        photosynthetic
                                                                                                                                                                                         porphyric herbicide resistance; triazine resistance; disease resistance;
                                                                                                                                                                                                                                                       abiotic stress tolerance; improved nutritional value; hygromycin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABK26344 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-106307/14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    consist of o-methyl modification, an LNA modification, two or more
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                respect to the genetic sequence to be altered and further comprises chemical modifications of the oligonucleotide. The chemical modifications
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               having a DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 6 A;
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83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ich comprises a single-stranded oligonucleotide The DNA domain has at least one mismatch with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 C; 5 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ç
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M
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Pred: No.
                                                                                                                                                                                                                                                                                                                genome altering oligonucleotide;
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                                                                                                                                      albino plant;
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                                                                                                                                                                                                                                                         primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local Similarity
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30-OCT-2000;
27-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               improved
modified
                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   respect to the genetic sequence to be altered and further comprises chemical modifications of the oligonucleotide. The chemical modification consist of o-methyl modification, an LNA modification, two or more
                                                                                                                                                                 Oligonucleotide
                                                 Homo sapiens
                                                                                                                                                                                                      21-FEB-2002
                                                                                                                                                                                                                                           ABF31424;
                                                                                                                                                                                                                                                                             ABF31424 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17
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                                                                                                                                                                                                  (first entry)
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2001US-0818875
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                                                                                                                                                               SEQ
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                                                                                                                                                           ID NO 131421 for detecting SNP TSC0032802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 C; 1 G; 6 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           80.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               to an oligonucleotide for targeted alteration ch comprises a single-stranded oligonucleotide
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Pred. No. 3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                designed to detect single nucleotide methylation status \boldsymbol{\cdot}
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Set of oligonucleotides,
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Pred. No. 6.8e
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                                                      This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                  Set of oligonucleotides, udesigned to detect single methylation status
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Pred. No. 6.8e+03;
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central nervous system, oligomers are also used

ABC00010-ABC99989,

used for detecting ABF000010-ABF99989,

cardiovascular and metabolic disorders.

gastrointestinal,

respiratory

cell type differentiation ABH00010-ABH99989 and

range of diseases including immune system,

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ABF7
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ABF76055 standard; DNA; 13
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                                                                                                                                                                                                                                   acid (PNA) oligomers for detecting single nucleotide polymorphisms (S and cytosine methylation status in chemically pretreated genomic DNA oligonucleotides are used for diagnosis and/or prognosis of cancer an
                                                                                                                                                                                                                                                                                                                                          Set of oligonucleotides, useful for diagnosis and designed to detect single nucleotide polymorphisms methylation status
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                                                                                                 Sequence
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Pred. No. 6.8e+03;
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                                                                                            RESULT 10
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28-NOV-1997;
17-DEC-1997;
                                                                                                                                                                                                                                                                                against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal uretritis, epidymitis, cervicitis, salpingitis, perihepatitis, bartholinitis; pneumopathy in breast feeding infants; and venereal lymphogranulomatosis. The polypeptides of the
                                                                                                                                                                                                                                                                                                                                                                                          PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs encode polypeptides (see AAZ36754-Y37949) which can be used as vaccines
           22-FEB-2002
                                      ABI25946
                                                                 ABI25946
                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 1597; 1755pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   bartholinitis;
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         (first entry)
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97FR-0015041.
97FR-0016034.
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69.2%;
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Pred. No. 6.
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peptide nucleic acid; cytosine methylation;
central nervous system; gastrointestinal; re
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID 325919;
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                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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                                                                                                                                                                                                                                                                                                                                                                      Sequence 12 BP; 2 A; 4 C; 1
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                                                                                                                                          Oligonucleotide primer SEQ ID NO 378713 for detecting SNP TSC0006271
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primer; ss;
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06-APR-2001; 2001WO-IB00713

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)
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            Set of oligonucleotides, useful for diagnosis and cell designed to detect single nucleotide polymorphisms and methylation status \,
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range of diseases including immune system, gastrolitestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989 and ABF00010-ABF998
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9; Conser
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                                                                                                                                                                                                                                                                                                                             ID 160398; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              system;
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81.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gastrointestinal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Berlin
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Pred. No. 1.1e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                     for diagnosis and cell typing, is
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cc a sample. The method comprises (1) generating cDNA from polyadenylated CC RNA employing at least 1 primer hybridizing to the polyadenylated RNA and CC comprising a 5' poly(C) or a 5' poly(G) flank; (2) (either): (a) (comprising a 5' poly(C) or a 5' poly(G) flank; (2) (either): (a) comprising a 5' poly(C) flank; (2) (either): (a) primer(s) comprising a 5' poly(C) flank was/cc (G) tail when in step (1) primer(s) comprising a 5' poly(C) flank was/cc employed or a poly(C) tail when in step (1) primer(s) comprising of the cc generated cDNA with a poly(G) tail when in step (1) primer(s) comprising of the cc generated cDNA with a poly(G) tail when in step (1) primer(s) comprising of the cc generated cDNA with a poly(G) tail when in step (1) primer(s) comprising of cc generated cDNA with a poly(G) flank was/were employed or a poly(C) tail when in step (1) cc primer(s) comprising a 5' poly(G) flank was/were employed using an RNA-cc ligase, irrespective of the presence or absence of surplus primer(s) and cc phylidizing to the tails generated in step (2ai) or (2b). The amplified cc plank obtained may be used for in vitro and/or in vivo expression and cc preparation of mRNA transcripts (which may then be used in hybridization con account contains the primer con and cc preparation of mRNA transcripts (which may then be used in hybridization con account contains contai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                밁
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AAI66878/c
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Best Local
                                          assays (comprising hybridization to oligonucleotide arrays, ci
and/or PNA arrays) and/or interaction assays (comprising inter
with carbohydrates, lectins, ribozymes, proteins, peptides, ar
and/or aptamers)) and for sequence specific PCR, cDNA cloning,
subtractive hybridization cloning and/or expression cloning. S
AAI66876-890 represent primers used in the method of the inver-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A method for the amplification of mRNA from a sample, e.g. for preparation of in vitro surrogates for pathologically modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-MAR-2000; 2000EP-0106450
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   provides a novel method for the amplification
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81.8%;
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Mismatches 0
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Conservative

73.3%; 81.8%;

Score 11; DB 22; Pred. No. 1.2e+04; Mismatches

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RESULT 15
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                                                                                                                                                                                                                                         The present invention relates to the use of an inhibitor of the interferon gamma (IFN-gamma) signalling pathway for the preparation of a pharmaceutical composition for the treatment or prevention of restenosis. The inhibitor is useful for the preparation of a pharmaceutical composition useful in treating or preventing restenosis which include restenosis of coronary arteries, carotid arteries, which include restenosis of coronary vein bypass, arterial bypass, and/or venous bypass; and restenotic modification, where prevention and/or venous bypass; and restenotic modification and the restenosis or restenotic modification is done before, during and/or after ballon angioplasty and/or stent implantation and the restenosis or restenotic modification is in-stent restenosis; and for treating and preventing restenosis in a subject preferably human. The present sequence is a desmin PCR primer which is used in aberrant gene expression in human restenotic tissue used in the exemplification of the invention.
                                                                                             Query Match
Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Use of an inhibitor of the interferon-gamma signaling pathway for preparation of a pharmaceutical composition useful in the treatment or prevention of restenosis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200170953-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAD19670 standard; DNA; 20 BP
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                                                                                                                                                                                                  Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 5; Page 55; 151pp; English.
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5 AACGUGAAGAU 15
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13 AACGTGAAGAT 3
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                                                                                                                    73.3%;
                                                                                                                                                                                             6 C; 5 G; 5 T; 0 other;
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Pred. No. 1.2e+04;
Pred. No. 1.2e+04;
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Search completed: May 22, Job time: 148.364 secs 2003, 19:04:24

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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SUMMARIES	
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Result No.	Score	Query Match Length DB	ength DB	Query Match Length DB ID
<u>,</u>	10.2	68.0	19 1	AZ759898
c 2	8.8	58.7	19 1	AZ465132
(L)	8.6	57.3	19 1	AZ603744
C 4	8.6	57.3	20 9	AU254575
· U	8.4	56.0	20 1	AZ501985
6	8.2	54.7	19 1	C21102

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33 HUMOUCC31 1154 AUO61154 38918 1M0397J 72819 1M0584F 33477 2M0025D	1147 AU011147 41695 1M0504E2 07038 2M0069C0 07038 2M0129E0 035078 2M0129E0 035078 1MMGS00033	52631 1M0269F1 76391 1M0295F1 76391 5009-0-2 97047 5009-0-2 96805 1M0333G2 9934 ti63c04.x	180710 10006200 18837 10011861 14372 10024461 17223 10024461 17247 10024461 10096 10035482 10106 10035482 10106 1003910 10106 1003910	6629 IM0224 8282 IM0535 8282 IM0536 3624 IM0059 13235 IM0059 1326 IM0080 5531 IM0080 5531 IM0080 5572 IM0080

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	JOURNAL COMMENT	TITLE	REFERENCE	ACCESSION VERSION KEYWORDS SOURCE	RESULT 1 AZ759898 LOCUS DEFINITION
University of Utah Genome Center University of Utah Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT 84112, USA Tel: 801 585 5606 Fax: 801 585 7177 Email: ddunn@genetics.utah.edu Insert Length: 10000 Std Error: 0.00	plasmid inserts Unpublished (2000) Contact: Robert B. Weiss	Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 19) Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,	clone UUGCIM0553A08 F, DNA sequence. A2759898 A2759898.1 GI:12867157 GSS. House mouse.	AZ759898 19 bp DNA linear GSS 16-FEB-2001 1M0553A08F Mouse 10kb plasmid UUGC1M library Mus musculus genomic

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                    Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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                                                                                                                                   University of Utah
University of Utah
                                                                                                                                                                                                     Unpublished (2000)
                                          Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                  Contact: Robert B.
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ddunn@genetics.utah.edu
Length: 10000 Std Error: 0.00
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                                                                                                              Biomedical Polymers Research Bldg.,
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/strain="C57BL/6J"
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Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
                                                                                                                                      University of Utah
                                                                                                                                                                                    Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                                     ,M., Rose,M., Rose,R.,
and Wright,D.,Weiss,R.,
Mouse whole genome sca
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Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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                                                                                                                Biomedical Polymers Research Bldg.,
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/note="Vector: PWD42nv; Purified genomic DNA from M.
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/strain≈"C57BL/6J"
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                                                                                                                                                                 Contact: Kikuya Kato
Graduate School of Biological Sciences
Nara Institute of Science and Technology
8916-5 Takayama, Ikoma, Nara 630-0101, Jt
Tel: 81-743-72-5581
                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AU254575 20 bp mRNA linear E
AU254575 3'-directed mouse cDNA library Mus musculus
                                                                                                                                                                                                                                                                                           Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                     Kato, K. and Matoba, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BED0002532 3', mRNA sequence.
                                                                                              URL:http://love2.aist-nara.ac.jp/BED/index.html.
                                                                                                                                                                                                                                                                                                               Generation of expressed sequence tags from mouse brain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AU254575.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                    Email: kkato@bs.aist-nara.ac.jp,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 house mouse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor ollgonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNBA2 (91473114)ghAF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

2 c 5 g 3 t
/organism="Mus musculus"
/db_xref="taxon:10090"
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                                                                          location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (http://www.jax.org/resources/documents/dnares/). The DNA
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/clone="UUGC1M0423E15"
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/strain="C57BL/6J"
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60.0%;
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Pred. No. 5.7e+05;
2; Mismatches 4;
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| | | | :||||:
16 AAAAAGCCTGAAGAT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0341 row: K column: 05
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            University of Utah Genome Center University of Utah Rm. 308, Biomedical Polymers Res
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clone UUGC1M0341K05 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       High quality sequence stop: 20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (2000)
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M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed.
                                                                                                                                                                         (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant vellocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The capacitact of the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                     /note="Vector: PWD42nv; Purified genomic DNA from usculus C57BL/6J (male) was obtained from the
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/tissue_type="brain"
/note="Vector: pGEM-T-easy"
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                                                                                                                                                                                                                                                                                                                                                              /note="Vector:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                               /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                                                                                                                                                                                                                                                                                                             /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone="UUGC1M0341K05"
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Pred. No. 5:9e+05;
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                                   3 AUAACGUGAAGAU 15
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C21102.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1-3, Yamada-oka, Suita, Osaka Pref. 565, Japan Tel: 06-877-511.(ex.3315)
Email: kousakuéincb.osaka-u.ac.jp
Human Gene Signature, 3'-directed cDNA sequence. We are not submitting the same cDNA sequence redundantly to DDBJ since 1993. For the abundance information of clones with this sequence in this library and as well as in other 3'-directed libraries, see'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C21102 19 bp mRNA linear I HUMGS0002625 Human adult (K.Okubo) Homo sapiens cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           http://www.imcb.osaka-u.ac.jp/bodymap'. The sequences of the clones represented by this GS sequences is also found there.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Okubo, K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                            Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Osaka University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BodyMap; human gene expression database
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1 (bases 1 to 19)
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a 4 c 3 g 6 t
                                                                                                                                                                                                                                                                                            /note="Organ: blood; Vector: 1-gt-11; Site_1: Eco-RI; Monocytes were prepared from blood by ficol1-hypaque, percol1 and T cel1 rosetting purification steps (purity: 96 %). mRNA was prepared from activated monocytes from a patient with rheumatoid arthritis. mRNA was reverse
                                                                                                                                                                                                                                    transcribed with MuLV. Using Eco-RI linkers cDNA was cloned into 1-gt-11 vector arms. The cDNA library was screened by differential hybidization using radioacti
                                                                                                                                                                             monocytes.
                                                                                                                                                                                                                  marked ss-cDNA from activated and
                                                                                                                                                                                                                                                                                                                                                                                               /clone_lib="Human adult (K.Okubo)"
/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
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70.0%;
                                                                                              54.7%;
                                                                          Score 8.2; DB 14;
Pred. No. 9.5e+05;
3; Mismatches 3;
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Pred. No. 7.6e+05;
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RESULT 7 AZ436629

RESULT 8 AZ658282/c

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COMMENT
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                                                                                                                                                                                               BASE COUNT
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                                                                                      Matches
                                                                                                           Query Match
Best Local
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                                      2 GAUAACGUGAAGA 14
GAGAACTTGATGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhaus and Wright,D.,Weiss,R.
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1M0224019F Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0224019 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Insert Length: 10000 Std Error: Plate: 0224 row: O column: 19 see primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        University of Utah Genome University of Utah Rm. 308, Biomedical Polyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus
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AZ436629
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      plasmid inserts
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1 (bases 1 to 19)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AZ436629.1 GI:10560642
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: ddunn@genetics.utah.edu
                                                                                      9;
                                                                                                           Similarity
                                                                                      Conservative
                                                                                                                                                                                                                                                                                               was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114)gblAF129072 l), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and provided the sheared adaptored mouse DNA was annealed to be sheared.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
                                                                                                                                                                                                                                      purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically competent E. coli XL10-Gold (Stratagene) cells
                                                                                                                                                                                             and selected for ampicillin resistance.

1 c 8 g 5 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note-"Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0224019"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /lab_host="E. Coli strain XL10-Gold, T1-resistant,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="Mouse 10kb plasmid UUGClM library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ocation/Qualifiers
  19
                                                                                                           54.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Weiss
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                                                                                      Score 8.2; DB 17;
Pred. No. 9.5e+05;
1; Mismatches 3;
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                                                                                                                                                                                                                               (Stratagene) cells
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RESULT 9
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DEFINITION
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Best Local
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                                                                                         ATATCTTGAATAT 5
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Insert Length: 10000 Std Err
Plate: 0535 row: C column: 0
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Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: 801 585 5606
Fax: 801 585 7177
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1M0535C02F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0535C02 F, DNA sequence.
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University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (2000)
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                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                     adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gil4732114|gblAF129072.1), a copy-number inducible derivative of plasmid RI. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in holar excess. The
                                                                                                                                                                                                                                                                                                  adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

4 c 2 g 5 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       musculus C57BL/6J (male) was obtained from the Jackson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="Mouse 10kb plasmid UUGCIM library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:10090"
/clone="UUGC1M0535C02"
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/strain="C57BL/6J"
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'note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'sex="Male"
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                                                                                                                                                                                                      Score 8.2; DB 17; Pred. No. 9.5e+05;
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13
                                                                                                                       Local Similarity
                                           1 AGAUAACGUGAAG 13
AGATAATGAGCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.
Plate: 0264 row: A column: 22
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   84112, USA
Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert B. University of Utah University of Utah
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2M0264A22R Mouse 10kb plasmid UUGC2M library Mus musculus genomic
clone_UUGC2M0264A22 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., ISlam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly, ROSE,M., ROSE,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mammalia; Eutheria;
1 (bases 1 to 19)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                      polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated
                                                                                                                                                                                                                with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance. **

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4
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Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="UUGC2M0264A22"
/clone_lib="Mouse 10kb plasmid UUGC2M library"
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/strain="C57BL/6J"
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Pred. No. 9.5e+05;
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                                          1 AGAUAACGUGAAG 13
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Insert Length: 10000 Std Erro
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1M0378J08F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0378J08 F, DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished
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                                                                                                                                                                                                                                                                   10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                              (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                    adaptored vector DNA, and transformed into chemically-competent E. coli XLLO-Gold (Strand selected for ampicillin resistance."

5 c 3 g 4 t
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/clone="UUGC1M0378J08"
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/strain="C57BL/6J"
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'note="Vector: PWD42nv; Purified genomic DNA from M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone_lib="Mouse 10kb plasmid UUGCIM library"
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Rodentia;
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                                                                                                    Score 8.2; DB 17; Pred. No. 9.8e+05;
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RESULT 11 AZ331326/c

RESULT 12 AZ345449/c

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GGTAAGGTGAA 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0059 row: B column: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mammalia; Eutheria;
1 (bases 1 to 19)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tel: 801 585 5606 Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly, Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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                                                                                                                                  Similarity
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                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                              (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pMB42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
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                                                                                                                                                                                                                                        adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Laboratory Mouse DNA Resource
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/strain="C57BL/6J"
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Rodentia;
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                                                                                                      Score 7.8; DB 17;
Pred. No. 1.6e+06;
7. Mismatches 2;
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Sciurognathi; Muridae; Murinae; Mus
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GGTAAGGTGAA 3
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Insert Length: 10000 Std Erro
Plate: 0080 row: I column: 08
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Fax: 801 585 7177
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University of Utah
Rm. 308, Blomedical Polymers Research Bldg., 20 S.
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AZ345449.1 GI:10424686
GSS
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and Wright,D.,Weiss,R.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                         Similarity 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              house mouse.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
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/clone="UUGC1M0080I08"
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/strain="C57BL/6J"
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                                                                                                               52.0%;
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                                                                                         2;
                                                                                                          Score 7.8; DB 17;
Pred. No. 1.6e+06;
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RESULT 13 AZ345511/c

RESULT 14 AZ345536/c

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Insert Length: 10000 Std Error:
Plate: 0080 row: J column: 01
Seg primer: CGTTGTAAAACGACGGCCAGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus
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1M0080J01F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0080J01 F, DNA sequence.
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Location/Qualifiers
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Fax: 801 585 7177
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                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

9 c 0 g 6 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    musculus C57BL/6J (male) was of Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note-"Vector: PWD42nv; Purified genomic DNA from M.
nusculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sex="
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="Mouse 10kb plasmid UUGC1M library"
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/clone="UUGC1M0080J01"
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                                                                                                                                      Score 7.8; DB 17;
Pred. No. 1.6e+06;
2; Mismatches 2
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Insert Length: 10000 Std Erro
Plate: 0080 row: O column: 00
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AZ345536
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Location/Qualifiers
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Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2000)
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                           adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."

9 c 0 g 6 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
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/strain="C57BL/6J"
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                                                                                                                                                                                          52.0%;
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Rodentia;
                                                                                                                                                                 Score 7.8; DB 17;
Pred. No. 1.6e+06;
2; Mismatches 2;
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Sciurognathi; Muridae; Murinae; Mus
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GAUAACGUGAA 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0080 row: J column: 17
Seg primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               84112,
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Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               High quality sequence stop: 19.
Location/Qualifiers
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Fax: 801 585 7177
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University of Utah
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Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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l (bases 1 to 19)
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                                             Conservative
                                                                                                                                                                                                                                                 was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114)gblAF129072 l), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
                                                                                                                                                                             purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for amnicillin resistance "
                                                                                                                                                  and selected for ampicillin resistance. 9 c 0 g 6 t
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0080J17"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /lab_host="E. Coli strain XL10-Gold, T1-resistant, F
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Laboratory Mouse DNA Resource
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Rodentia;
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
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4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
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Listing first 45 summaries
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   DB
US-09-199-865-4
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US-08-1812-968A-447
US-08-78-306-1325
US-09-66-1325
US-09-780-143-70
US-09-561-497-82
US-09-844-634-63
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Sequence 70, Appl
Sequence 63, Appli
Sequence 5, Appli
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CURRENT APPLICATION NUMBER: US/09/780,175

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SEQUENCE CHARACTERISTICS:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                         TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                               FILING DATE: 13-JANU PRIOR APPLICATION DATA:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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TITLE OF INVENTION:
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ZIP: 90071-2066
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REGISTRATION NUMBER: 3:
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VENTION: METHOD AND REAGENT FOR VENTION: INHIBITING HEPATITIS C
VENTION: VIRUS REPLICATION
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Pred. No. 1
                                                Mismatches
                                                          3.8e+03;
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                                                                         DB 1;
                                                                        Length 15;
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                                               Indels .
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                                               Gaps
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: Sequence 447, Application US/09064156A

: Patent No. 6132966
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                                                                                                                                                                                                                                                                                                            Matches
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                                         APPLICANT: Draper, Kenn
TITLE OF INVENTION: METI
TITLE OF INVENTION: VIRI
TITLE OF INVENTION: VIRI
NUMBER OF SEQUENCES: 49:
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/18
FILING DATE: January 13,
APPLICATION NUMBER: 07/88
FILING DATE: May 14, 1992
ATTURNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Word Perfect 5
CURRENT APPLICATION DATA:
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TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (213) 489-1600
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TELEX: 67-3510
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STATE: California
                               ADDRESSEE:
                                                                                                                                                                                                                                                                                                           Local Similarity 66.7 es 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Warburg, Richard J
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                            1 AGAUAACGUGAAGAU 15
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                                                                                                                                                                                                                                                AGATAACGACAAGGT 1
                                                                                                                                       INFORMATION:
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90071-2066
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5869253
                                                                                                                                                                                                                                                                                                                                                                                                                nucleic acid
633 West Fifth
Suite 4700
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633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                     is: single
linear
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                              Lyon & Lyon
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                                                                                     Kenneth G.
METHOD AND REAGENT FOR
INHIBITING HEPATITIS C
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METHOD AND REAGENT FOR
INHIBITING HEPATITIS C
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66.7%;
                                                                            VIRUS REPLICATION
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ber 26, 1996
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               Street
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Pred. No. 3
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Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                        atent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 234/083
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15 AGATAACGACAAGGT 1
                                                                                            COUNTRY:
                                                                                                                        CITY: Abbott Park
                                                                                                                                        STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: linear
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REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE: December
                                                                                                                                                        ADDRESSEE:
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APPLICATION NUMBER: 08/182,968
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                        INFORMATION:
                                                                               60064
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90071-2066
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                                                                                                                                   100 Abbott Laboratories
                                                                                            USA
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                                                                                                                                                                                                                               Dawson, G.
                                                                                                                                                                                                                                              Mushahwar, 1
Chalmers, M
                                                                                                                                                                                                                                                                               Desai,
                                                                                                                                                                                                                                                                                          Erker,
                                                                                                                                                                                                                                                                                                                          Birkenmeyer, L.
                                                                                                                                                                                                                                                                                                                                          Simons, J. N.
PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                          Muerhoff,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IBM Compatible
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JMBER: 07/882,888
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                                                                                                                                                                                                 Genotypes
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66.7%;
                                                                                                                                                                                                              Detection of Hepatitis GB Virus
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ber 26, 1996
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Pred. No. 3
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                                                                                                                                                                    US-08-758-306-1325/c
                                                                                                                                                                                     RESULT
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US-09-780-175-50/c
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                                                                                                                Patent No. 580, ...
                                                                                                                                         Sequence 1325, Application US/08758306 Patent No. 5807743
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 50, Application US/09780175 Patent No. 6440738
                                                                                                                                                                                                                                                                                        Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: RTS-0164
CURRENT APPLICATION NUMBER: US/09/780,175
CURRENT FILING DATE: 2001-02-08
                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
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NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
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                                                                      TITLE OF INVENTION:
                                                                                                  APPLICANT:
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                                                                                                                                                                                                                                19 ACGTGAAGAT 10
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AGAGAAGGTTAAGAT 18
                                                                                                                                                                                                                                                                                      Similarity
8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Susan M. Freier
              SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jacqueline Wyatt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Robert McKay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       708-938-2623
                                                                                              McSwiggen,
                                                                                                             Stinchcomb, Dan T.
                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ANTISENSE MODULATION OF CASEIN KINASE 2-BETA EXPRESSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           68.0%;
                       GAMMA-CHAIN EXPRESSION
                                                                                                                                                                                                                                                                                                     80.0%;
                                                     ASSOCIATED WITH
                                                                                  n, James A.
METHOD AND REAGENT FOR THE
                                                                    TREATMENT OF DISEASES
                                         INTERLEUKIN-2
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                                                                                                                                                                                                                                                                                                    Score 10;
Pred. No.
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Pred. No. 3
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COUNTRY:

Los Angeles California STREET:

E: Lyon & Lyon 633 West Fifth Suite 4700

Street

ADDRESSEE:

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                                                                                                                      ; OTHER INFORMATION: Antisense Oligonucleotide US-09-205-143-70
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                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/205,143
CURRENT FILING DATE: 1998-12-03
NUMBER OF SEQ ID NOS: 87
SEQ ID NO 70
LENGTH: 18
                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                    Sequence 70, Application US/09205143 Patent No. 6107091
                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                         APPLICANT: Lex M. Cowsert TITLE OF INVENTION: ANTISENSE MODULATION OF G-ALPHA-16 EXPRESSION FILE REFERENCE: RTS-0032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                   FEATURE:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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APPLICATION NUMBER:
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REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US
FILING DATE: December
                                                                                                                                                                                                                                                                                                                                                                                                                               13 AGATAAGGGGAAG 1
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TOPOLOGY: lin
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18 GAAAAGGTGAAGA 6
                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 AGAUAACGUGAAG 13
                            2 GAUAACGUGAAGA 14
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                                                           Conservative
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76.9<del>8</del>;
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                                                                       76.9%;
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er 3, 1996
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Pred. No.
                                                                       Score 9.8;
Pred. No. 6.
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                                                           Mismatches
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                                                                    .5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-09-561-497-82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-561-497-82
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SEQ ID NO 82
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 174
SEQ ID NO 63
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 82, Appli
Patent No. 6372433
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Best Local Similarity
Matches 9; Conser
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Best Local Similarity
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                                                                                                                                                                                  Sequence 5, Application US/08180209B Patent No. 5593877
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/844,634
CURRENT FILING DATE: 2001-04-27
NUMBER OF SEQ ID NOS: 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/561,497 CURRENT FILING DATE: 2000-04-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Andrew T. Watt TITLE OF INVENTION: ANTISENSE MODULATION OF TUMOR NECROSIS FACTOR RECEPTOR 2 EXPREFILE REFERENCE: RTS-0216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Brenda F. Baker
APPLICANT: C. Frank Bennett
                                                                                                                                                                    GENERAL INFORMATION:
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                                                                     APPLICANT: King, Te-Piao
TITLE OF INVENTION: CLOWIN
TITLE OF INVENTION: VESPID
TITLE OF INVENTION: HYALUR
TITLE OF INVENTION: THEREO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
                                   CORRESPONDENCE ADDRESS:
                                                     NUMBER OF SEQUENCES:
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2 AGCTAACTTGAAG
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Klauber & Jackson
Hackensack Avenue
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69.2%;
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84.6%;
                                                                                                             CLONING AND RECOMBINANT PRODUCTION OF VESPID VENOM ENZYMES, SUCH AS PHOSPHO:
                                                                                         HYALURONIDASE, AND IMMUNOLOGICAL THERAPIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 9.8; DB 4;
Pred. No. 6.5e+03
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Pred. No. 6.5e+03;
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COUNTRY:

USA

Hackensack : New Jersey

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US-08-385-745-5
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: King, Te Piao
TITLE OF INVENTION: Clonin
TITLE OF INVENTION: Vespid
TITLE OF INVENTION: Based
NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENERAL INFORMATION:
                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: DNA
                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
NAME: Misrock, S. Leslie
                                                                                                                                                                                                                                                                                                                                                            ORRESPONDENCE ADDRESS:
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APPLICATION NUMBER: US 08/031,400

FILING DATE: 11-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: sin
TOPOLOGY: linear
                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                   STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Match 64.0%; Local Similarity 50.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/0 FILING DATE: 11-JAN-1994
                                                                                                                                                                                                                                                    COUNTRY: U.S.A. ZIP: 10036-2711
                                                                                                                                                                                                                                                                                                                        STREET:
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REFERENCE/DOCKET NUMBER: 600-1-074 CIP
                                   FILING DATE:
                                                                                       CLASSIFICATION: 435
                                                                                                          FILING DATE:
                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                        CITY: New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Jackson Esq., David A. REGISTRATION NUMBER: 26,742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
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1155 Avenue of the Americas
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                                                    US/08/031,400
                                                                                                                       US/08/385,745
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Pred. No.
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Best Local Similarity
Matches 7; Conserve
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; MOLECULE TYPE:
US-08-385-745-5
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                                               INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base ---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 5, Application US/08485388 Patent No. 6270763
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                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: UFILING DATE: 07-JUN-1
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: 212 790-9090
TELEFAX: 212 869-8864/9741
TELEX: 66141 PENNIE
                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 08/38
FILING DATE: 08-FEB-1995.
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MOLECULE TYPE:
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                                                                                                                                                     RELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 11-MAR CLASSIFICATION: 435
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                STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                        NAME: Jackson Esq., David A. REGISTRATION NUMBER: 26,742 REFERENCE/DOCKET NUMBER: 600
                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM:
SOFTWARE: PatentI
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: New Jersey
                                         nucleic acid
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Pred. No. 8
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Best Local Similarity
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US-08-474-853-5
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                                                                      <u>В</u>
                                                                                                      δÃ
                                                                                                                                                                                                           ; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: DNA
US-08-474-853-5
Search completed: May 22, 2003, 20:18:23 Job time: 35.2727 secs
                                                                                                                                       Query Match
Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 5, Application US/08474853 Patent No. 6287559
                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/474,853
FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/180,209
FILING DATE: 11-JAN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/031,400
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REGISTRATION NUMBER: 26,742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION: APPLICANT: King,
                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 26,742

REFERENCE/DOCKET NUMBER: 600-1-074 CIPB
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201 487-5800
TELEPAX: 201 343-1684
TELEX: 133521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORRESPONDENCE ADDRESS:
ADDRESSEE: Klauber & Jackson
                                                                                                                                                                                                                                                                                                                                      TELEFAX: 201
TELEFAX: 133521
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Hackensack
STATE: New Jersey
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: UZIP: 07601
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1 GAYACNGTNAARAT 14
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                                                                                                                                         Conservative
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Pred. No.
                                                                                                                                         Score 9.6; DB 4;
Pred. No. 8.3e+03;
4; Mismatches 3
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GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

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Minimum DB
Maximum DB
                                                                                                                                                                                                                                                                                       Result
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Maximum Match 1008
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Run on:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    seq
      length: 0 length: 20
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Match
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Gapop 10.0 , Gapext 1.0
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/cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
/cm2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
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/cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
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/cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
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/cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
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                                     US-10-000-773A-6

US-09-776-474-739

US-09-776-474-739

US-09-918-187-64

US-09-918-187-64

US-09-800-631-141

US-09-730-289B-596

US-09-78-582-29

US-09-969-373-1862
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US-09-730-289B-606
US-09-730-289B-607
US-10-213-329-4
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                US-09-917-963-42
US-09-953-611-56
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US-09-504-231A-469
US-09-274-553D-469
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                                    Sequence 97, Appl
Sequence 606, App
Sequence 607, Appli
Sequence 4, Appli
Sequence 469, App
Sequence 469, Appl
Sequence 739, App
Sequence 739, Appl
Sequence 64, Appl
Sequence 64, Appl
Sequence 141, Appl
Sequence 29, Appl
Sequence 29, Appl
Sequence 29, Appl
Sequence 4497, Appl
Sequence 4497, Appl
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Sequence 42, Appl
Sequence 56, Appl
Sequence 112, App
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ALIGNMENTS

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RESULT 2
US-09-730-289B-606/c
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APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Breaker, Ronald
APPLICANT: Beigelman, Leo
                                                                                                                                                                                           Sequence 606, Application US/09730289B Publication No. US20030050259A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: MBHB00-884-H (500/001)
CURRENT APPLICATION NUMBER: US/09/780,929
CURRENT FILING DATE: 2001-02-08
PRIOR APPLICATION NUMBER: US 60/181,360
PRIOR FILING DATE: 2000-02-08
NUMBER OF SEQ ID NOS: 126
APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Blatt, Larry APPLICANT: MCSwidgen, Jim TITLE OF INVENTION: Method and Reagent for the APPLICANT of Invention of the APPLICANT APPLICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: RNA
ORGANISM: Artificial Sequence
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and Reagent for Treatment of Cardiac Disease
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; TYPE: RNA; Homo sapiens US-09-730-289B-606
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Best Local Similarity
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                                                                                                                                                                                                                                                                                             Sequence 4, Application US/10213329 Publication No. US20030083465A1
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for Treatment of Cardiac Disease
FILE REFERENCE: Method and Reagent for Treatment of Cardiac Disease
CURRENT APPLICATION NUMBER: US/09/730,289B
CURRENT FILING DATE: 2000-12-05
PRIOR APPLICATION NUMBER: US 60/169,100
PRIOR FILING DATE: 1999-12-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 3897
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/730,289B CURRENT FILING DATE: 2000-12-05
                                                                                                                                                                                             APPLICANT: Zimrin, Ann B.
APPLICANT: Maciag, Thomas
APPLICANT: Wong, Michael K.K.
APPLICANT: Pepper, Michael S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Ribozyme Pharmaceuticals, Inc.
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                                                     PRIOR APPLICATION NUMBER: US/09/199,865 PRIOR FILING DATE: 1998-11-25
                                                                                         CURRENT APPLICATION NUMBER: US/10/213,329
CURRENT FILING DATE: 2002-08-06
                                                                                                                                            TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC METHODS AND COMPOSITIONS TITLE OF INVENTION: BASED ON JAGGED/NOTCH PROTEINS AND NUCLEIC ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: MBHB00-864-A (400/006)
                                                                                                                              FILE REFERENCE: 0036-101
                                   PRIOR APPLICATION NUMBER:
                                                                                                                                                                                  APPLICANT: Montesano, Roberto
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17 AGATAACATGAGGA 4
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PCT/US97/09407
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Pred. No. 1
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Pred. No. 1
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LENGTH: 18
TYPE: DNA
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LENGTH: 19
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APPLICANT: Effertz, Roger J.
APPLICANT: Hauge, Brian M.
                                                                                                                                                                                                                                                  Sequence 469, Application US/09504231A Patent No. US20020013458A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local 9
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2001-01-13
PRIOR APPLICATION NUMBER: US 09/855,768
PRIOR FILING DATE: 2001-05-15
                 CURRENT APPLICATION NUMBER: US/09/504,231A CURRENT ETILING DATE: 2000-02-15 PRIOR APPLICATION NUMBER: 09/274,553 PRIOR FILING DATE: 1999-03-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/969,373
CURRENT FILING DATE: 2001-10-02
PRIOR APPLICATION NUMBER: US 09/754,853
PRIOR EILING DATE: 2001-01-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS:
                                                                                             TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF TITLE OF INVENTION: HEPATITIS C VIRUS INFECTION FILE REFERENCE: rpl 247/282
                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Hauge, Brian M.
TITLE OF INVENTION: Soybean SSRs and Methods of Genotyping
FILE REFERENCE: 38-10(52679)A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Ver. 2.0.
                                                                                                                                                                                                                                APPLICANT: Blatt,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 4593
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 09/760,427
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 1997-05-30
                                                                                                                                                       APPLICANT:
                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: cDNA OTHER INFORMATION: amplification 5'-primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
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                                                                                                                                                     Macejak, Dennis
                                                                                                                                                                        Pavco, Pamela
                                                                                                                                                                                          Roberts, Beth
                                                                                                                                                                                                          McSwiggen, James
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71.4%;
09/257,608
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Pred. No. 2
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RESULT 8
US-10-000-773A-6
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                                                                                 Sequence 6, Application US/10000773A Publication No. US20030069195A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                             Query Match
Best Local :
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 469
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Best Local Similarity
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                                                  APPLICANT: Farrar, APPLICANT: Humphr
APPLICANT: Millington-Ward, Sophia
                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                      LENGTH: 15
TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/100,842
PRIOR FILING DATE: 1998-09-18
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CURRENT APPLICATION NUMBER: US/09/274,553D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: MACEJAK, Dennis
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF DISEASES
TITLE OF INVENTION: HEPATITIS C VIRUS INFECTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/083,217 PRIOR FILING DATE: 1998-04-27
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PRIOR FILING DATE: 1998-09-18
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15 AGATAACGACAAGGT 1
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AGATAACGACAAGGT 1
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                                Humphries, Peter
Kenna, Paul
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                                                                 Gwenyth Jane
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66.7%;
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66.7%;
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Pred. No. 2
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                                                                                                                                                                                                                                                                     2.7e+04;
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                                                                                                                                                                                                                                                                                                                               Nucleic Acid Target
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RESULT 10
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                                                                                                           Query Match
Best Local Similarity
Matches 11; Conserv
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Best Local Similarity 66.7
Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 739,
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                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/776 474
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: US 60/179,983
PRIOR FILING DATE: 2000-03-02
NUMBER OF SEQ ID NOS: 2992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Jarvis, Thale
APPLICANT: Booher, Robert
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TITLE OF INVENTION: Method and Reagent for
TITLE OF INVENTION: Enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: GB 9604449.0 PRIOR FILING DATE: 1996-03-01 NUMBER OF SEQ ID NOS: 34
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CURRENT FILING DATE: 2001-11-30
PRIOR APPLICATION NUMBER: US 09/142,125
PRIOR FILING DATE: 1999-04-12
PRIOR APPLICATION NUMBER: PCT/GB97/00574
PRIOR FILING DATE: 1997-03-03
                                                                                                                                                                                                                           TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: MBHB00-955-A (400/008)
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TYPE: DNA
ORGANISM: Artificial Sequence
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                                               2 AGAAAAUGUGAAG
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                                                                                                                    Conservative
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                                                                                                                                   65.3%;
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Pred. No. 4.
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Pred. No. 2
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APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Jarvis, Thale APPLICANT: Booher, Robert

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                                                                                                                                                                                                                                      US-09-800-631-141/c
                                                                                                                                                                                                                                                           RESULT 12
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Best Local S
Matches 11
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                  Sequence 141, Application US/09800631 Patent No. US20020082228A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 64
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FILE REFERENCE: MBHB00-955-A (400/008)
CURRENT APPLICATION NUMBER: US/09/776,474
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: US 60/179,983
PRIOR FILING DATE: 2000-03-02
NUMBER OF SEQ ID NOS: 2992
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CURRENT FILING DATE: 2001-07-30
NUMBER OF SEQ ID NOS: 80
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: ROSANNE M. CYOOKE
APPLICANT: MAIK J. GATAAM
TITLE OF INVENTION: ANTISENSE MODULATION OF STEAROYL-COA DESATURASE EXPRESSION
FILE REFERENCE: ISPH-0590
                                  CURRENT APPLICATION NUMBER: US/09/800,631
CURRENT FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: US/09/657,346
PRIOR FILING DATE: 2000-09-07
                                                                                                         APPLICANT: Hong Zhang
APPLICANT: Jacqueline Wyatt
TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: ISPH-0544
NUMBER OF SEQ ID NOS: 175
EQ ID NO 141
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                 4 ATATCCTGAAGAT 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Holman, Patricia
Fattaey, Ali
                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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61.5%;
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Pred. No. 4.4e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                            DB 9;
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                                                                                                                            BH3 INTERACTING DOMAIN DEATH AGONIST EXPI
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Best Local 9
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TYPE: DNA
                                    FEATURE:
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RESULT 13
US-09-730-2898-596
US-09-730-2898-596
; Sequence 596, Application US/09730289B
; Publication No. US20030050259A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-09-800-631-141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; ORGANISM: Homo sapiens US-09-730-289B-596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 14
US-09-878-582-29/c
: OTHER INFORMATION: Antisense Oligonucleotide
15-09-878-582-29
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Best Local Similarity 69.7
Matches 9; Conservative
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                                                                                                  NUMBER OF SEQ ID NOS: 51
SEQ ID NO 29
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No. US20020058638A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 29,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: McSwiggen, Jim TITLE OF INVENTION: Method and Reagent for Treatment of Cardiac Disease FILE REFERENCE: MBHB00-864-A (400/006) CURRENT APPLICATION NUMBER: US/09/730,289B CURRENT FILING DATE: 2000-12-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
                                                                                                                                                                                                                                        APPLICANT: ROBERT MCKAY
TITLE OF INVENTION: ANTISENSE MODULATION OF PTEN EXPRESSION
FILE REFERENCE: ISPH-0463
CURRENT APPLICATION NUMBER: US/09/878,582
CURRENT FILING DATE: 2001-06-11
PRIOR APPLICATION NUMBER: 09/577,902
PRIOR FILING DATE: 2000-05-24
PRIOR FILING DATE: 2000-05-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/169,100 PRIOR FILING DATE: 1999-12-06 NUMBER OF SEQ ID NOS: 3897
                                                                                                                                                               PRIOR APPLICATION NUMBER: PCT/US99/29594 PRIOR FILING.DATE: 1999-12-14
                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 09/358,381 PRIOR FILING DATE: 1999-07-21
                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Brett P. Monia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
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                                                             ORGANISM: Artificial Sequence
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69.2%;
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Pred. No. 7.3e+04
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Pred. No.
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RESULT 15

US-09-969-373-4497/c

Sequence 4497, Application US/09969373

Patent No. US20020133852A1

GENERAL INFORMATION:
APPLICANT: Effertz, Roger J.
APPLICANT: Hauge, Brian SSRs and Methods of Genotyping FILE REFERENCE: 38-10(52679)A

CURRENT APPLICATION NUMBER: US/09/969,373

CURRENT FILING DATE: 2001-10-02

PRIOR APPLICATION NUMBER: US 09/754,853

PRIOR FILING DATE: 2001-01-05

PRIOR FILING DATE: 2001-01-13

PRIOR APPLICATION NUMBER: US 09/760,427

PRIOR APPLICATION NUMBER: US 09/760,427

PRIOR APPLICATION NUMBER: US 09/760,427

PRIOR FILING DATE: 2001-01-13

PRIOR APPLICATION NUMBER: US 09/855,768

PRIOR APPLICATION NUMBER: US 09/855,768

PRIOR FILING DATE: 2001-05-15

NUMBER OF SEQ ID NOS: 4593

SEQ ID NO 4497

LENGTH: 18

Type: DNA
ORGANISM: Glycine max

US-09-969-373-4497
Search completed: May 22, 2003, 21:38:46 Job time: 72.9091 secs
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Best Local Similarity 72.7
Matches 8; Conservative
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                                                                                            2 GAUAACGUGAA 12
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|3 GATAACGTGCA 3
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17 AAGGTGAAGAT 7
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72.7%;
                                                                                                                                                                                  Score 9.4; DB 10; Length 18; Pred. No. 7.3e+04; 2; Mismatches 1; Indels
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Minimum DB
Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OM nucleic - nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     seq length: 0 seq length: 20
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Gapop 10.0 , Gapext 1.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24791104 segs, 12571243825 residues
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Maximum Match 100%
Listing first 45 summaries
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                    ptodata/1/pna/US101B_COMB.seq
                                     'ptodata/1/pna/US1099D_COMB.seq
'ptodata/1/pna/US109A_COMB.seq
'ptodata/1/pna/US100A_COMB.seq
'ptodata/1/pna/US100B_COMB.seq
                                                                                                                                                                                                                                                                                                                                                                                   Pending Nucleic Acid and/or Pending Amino Acid database searches now generate two sets of
                                                                                                                                                                                                                                                                                                                                                               results. These databases were split into to two parts to reduce the time needed to update the
                                                                                                                                                                                                                                                                                                                                         databases daily. The split freed up more machine time for processing searches.
                                                                                                                                                                                                                                                                                            Searches run against the Nucleic Acid Pending database produce two sets of results, with the
                                                                                                                                                                                                                                               Searches run against the Amino Acid Pending database produce two sets of results, with the
                                                                                                                                                                                                                           extensions, .rapm and .rapn
                                                                                                                                                                            The Pending database search results should not be left in the case because they
                                                                                                                                                    contain data that is confidential.
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/ptodata/1/pna/US
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      US-09-068-506-50

US-09-514-000-12603

US-09-514-000-15028

US-09-703-708-12172

US-09-703-708-12172

US-60-164-320-12172

US-60-184-3791-12172

PCT-US98-26935-217
                                                                             US-60-216-745-5964
PCT-US02-03159-22
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                                            Sequence 50, Appl
Sequence 12603, A
Sequence 15028, A
                                                                                                                   Sequence 606,
Sequence 607,
Sequence 4, A
Sequence 4, A
                                                                             Sequence 22, Appl
                                                                                                                                                         Sequence :
                                                                                     Sequence 4, Appl
Sequence 5964, A
                                      Sequence
                                                                                                          Sequence
                                                                                                       4, Appli
4, Appli
12, Appl
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RESULT 15
US-09-514-000-12603/c
; Sequence 12603, Application US/09514000
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US-09-068-506-50/c
; Sequence 50, Application US/09068506A
; GENERAL INFORMATION;
; APPLICANT: YASUE, Hirofumi
; APPLICANT: YOSHIMURA, Kumamoto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             밁
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                                                                                                                                                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
; OTHER INFORMATION: Primers
US-09-068-506-50
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PCT-US02-03159-22/c
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; OTHER INFORMATION: Antisense Oligonucleotide
PCT-US02-03159-22
                                                                                                                                                        Query Match 72.0%;
Best Local Similarity 64.3%;
Matches 9; Conservative
                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Ver.
SEQ ID NO 50
LENGTH: 20
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Best Local Similarity
Matches 9; Conserv
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LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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Best Local Similarity
Matches 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/068,506A CURRENT FILING DATE: 1998-07-10 NUMBER OF SEQ ID NOS: 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: PCT/US02/03159
CURRENT FILING DATE: 2002-01-31
NUMBER OF SEQ ID NOS: 154
                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: DIAGNOSIS OF DISEASES ASSOCIATED WITH CORONARY TITLE OF INVENTION: TWITCHING FILE REFERENCE: 0032-245P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: JACQUELINE WYATT
TITLE OF INVENTION: ANTISENSE MODULATION OF CASEIN KINASE 2-BETA EXPRESSION
FILE REFERENCE: RTSP-0270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
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14 GGTAACCTGAAGAT 1
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2 GATAACGTGCAAAT 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Susan M. Freier
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64.3%;
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                                                                                                                                                                      Score 10.8;
Pred. No. 1
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                                                                                                                                                      Mismatches
                                                                                                                                                                      1e+05;
                                                                                                                                                                                     DB 14;
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                                                                                                                                                                                  Length 20;
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Search completed: May 22, 2003, 21:22:03 Job time: 1731.64 secs
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                                                                                                                                                                                                         ; ORGANISM: Agrobacterium tumefaciens US-09-514-000-12603
                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/514,000
CURRENT FILING DATE: 2000-02-23
NUMBER OF SEQ ID NOS: 15034
SEQ ID NO 12603
.LENGTH: 20
TYPE: DNA
                                                                                                                                      Matches
                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
                                                                                                                                      Best Local Similarity Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15490)B
                                                                   20 GATGACCTGAAGAT 7
                                                                                                   2 GAUAACGUGAAGAU 15
                                                                                                                                      Conservative
                                                                                                                                                    72.0%;
                                                                                                                                                    Score 10.8;
Pred. No. 1
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le+05;
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GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

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Database :
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                 Score
    10.8
10.8
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110.4
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110.4
110.4
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Match
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    Pending_Patents_NA_New: *
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/cgn2_6/ptodata/2/pna/US60_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US60_NEW_COMB.seq2:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /ggn2_6/ptodata/2/pna/US06_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US07_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US08_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US09_NEW_COMB.seq:*
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/ptodata/2/pna/US09_NEW_COMB.seq3:*
/ptodata/2/pna/US10_NEW_COMB.seq:*
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US-10-302-027-46
US-10-302-027-106
US-10-31-109-21
US-09-846-903-24
US-09-846-903-24
US-10-310-188-80491
PCT-US03-03662-336
US-10-310-188-14089
PCT-US03-38806-82
US-10-317-277A-74
US-10-317-318-22900
US-10-318-23900
US-10-349-143-4165
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US-10-287-818-374
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73, Appl
148, App
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836, App
47798, A
                        469, App
22900, A
118, App
118, App
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336, App
14089, A
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24, Appl
80491, A
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106, App
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US-10-305-273-1246	US-10-305-273-723	US-60-453-420-94	US-10-385-163-94	US-10-089-177-1165	US-10-364-516-3268	US-10-364-516-3267	US-10-364-516-2146	US-10-364-516-2145	US-10-305-276A-1129	US-10-316-954-5747	US-10-316-954-4481	US-10-305-276-1129	US-10-287-787-13654	PCT-US02-28539-3268	PCT-US02-28539-3267	PCT-US02-28539-2146	PCT-US02-28539-2145	US-10-317-249-161	US-10-317-249-86	US-10-310-188-35003	US-10-298-994-212	US-10-298-994-110
Sequence 1246, Ap	-	Sequence 94, Appl	Sequence 94, Appl	Sequence 1165, Ap	Sequence 3268, Ap	Sequence 3267, Ap	Sequence 2146, Ap	`	•	`	Sequence 4481, Ap	`	**	Sequence 3268, Ap	•	•	'n	Sequence 161, App	Sequence 86, Appl	2	Sequence 212, App	Sequence 110, App

ALIGNMENTS

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; FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-10-316-389-88
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US-10-316-389-88/c
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Matches
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CURRENT FILING DATE: 2002-12-10
NUMBER OF SEQ ID NOS: 143
SEQ ID NO 88
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                     PRIOR APPLICATION NUMBER: 10/003,919
PRIOR ETILING DATE: 2001-12-06
NUMBER OF SEQ ID NOS: 87
SEQ ID NO 50
LENGTH: 20
                                                                                                                                                                                                                      Sequence 50, Application PC/TUS0238622
GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: Susan M. Freier
APPLICANT: Isis Pharmaceuticals, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 88, Application US/10316389 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                              APPLICANT: Susan M. Freier
APPLICANT: Isis Pharmaceuticals, Inc.
TITLE OF INVENTION: ANTISENSE MODULATION OF SHIP-1 EXPRESSION
FILE REFERENCE: RTSP-0447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Kenneth W. Dobie
TITLE OF INVENTION: ANTISENSE MODULATION OF ABCC2 EXPRESSION
FILE REFERENCE: RTS-0382
                                                                                                                                         CURRENT APPLICATION NUMBER: PCT/US02/38622 CURRENT FILING DATE: 2002-12-04
TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity hes 12; Conserv
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Pred. No. 1
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; LOCATION: (810101)...(810115)
; OTHER INFORMATION: Chromosome = 1 Strand = positive ConnectronObjectNumber = US-10-287-818-374
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Best Local Similarity
""" 9; Conserv:
RESULT 5
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CURRENT FILING DATE: 2002-11-05
NUMBER OF SEQ ID NOS: 1318
SOFTWARE: Proprietary
SEQ ID NO 374
                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 1318
SOFTWARE: Proprietary
SEQ ID NO 836
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Feldmann, Richard J.; Global Determi
TITLE OF INVENTION: Vibrio cholerae chromosome
                                                                                                                                  Query Match
Best Local :
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Best Local
                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: Jim Zegeer Law Offices - CURRENT APPLICATION NUMBER: US/10/287,818
CURRENT FILING DATE: 2002-11-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE: OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                          ORGANISM: Vibrio cholerae chromosome I, complete chromosome. FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                          LOCATION: (1812221)...(1812235)
OTHER INFORMATION: Chromosome = 1
                                                                                                                                                                                                                                                        TYPE: DNA
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                                                                                                                  l Similarity
9; Conserv
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                                                                                                                                73.3%;
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Pred. No.
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Pred. No. 1
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Pred. No. 2.9e+04;
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2.9e+04;
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RESULT 7
US-10-302-027-106
; Sequence 106, Application US/10302027
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-302-027-46/c

; Sequence 46, Application US/10302027
                                                  US-10-302-027-106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Homo sapiens US-10-310-188-47798
                                                                                                 NUMBER OF SEQ ID NOS: 135
SEQ ID NO 106
LENGTH: 20
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 135
SEQ ID NO 46
LENGTH: 20
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APPLICANT: RosettaGemonics
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Best Local
Query Match
Best Local Similarity
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CURRENT FILING DATE: 2002-12-19
NUMBER OF SEQ ID NOS: 86841
                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/302,027
CURRENT FILING DATE: 2002-11-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/302,027 CURRENT FILING DATE: 2002-11-21
                                                                                                                                                                                                                                     APPLICANT: Nicholas M. Dean APPLICANT: Kenneth W. Dobi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Nicholas M. Dean APPLICANT: Kenneth W. Dobi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL TITLE OF INVENTION: USES THEREOF
                                                                                                                                                                                                   FILE REFERENCE: PTS-0068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: PTS-0068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: MODULATION OF GANKYRIN EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: 47487
                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                ORGANISM: H. sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
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64.3%;
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Score 10.8; DB 9; Pred. No. 3.8e+04;
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Pred. No. 3.
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; OTHER INFORMATION:
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US-09-846-903-24
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                                                                                                                                                                                                                    SEQ ID NO 24
LENGTH: 17
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APPLICANT: Masucci, James D.
TITLE OF INVENTION: PLANT REGULATORY SEQUENCES FOR SELECTIVE CONTROL
TITLE OF INVENTION: OF GENE EXPRESSION
FILE REFERENCE: 38-21(15678)8 promoters
CURRENT APPLICATION NUMBER: US/09/846,903
CURRENT FILING DATE: 2001-04-26
PRIOR FILING DATE: 2000-05-01
PRIOR FILING DATE: 2000-05-01
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                                                                 Query Match
Best Local Similarity
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PRIOR FILING DATE: 2000-07-01
NUMBER OF SEQ ID NOS: 34
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PRIOR FILING DATE: 2001-07-02
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                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 98 SOFTWARE: PatentIn Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Conner, Timothy W.
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TYPE: DNA
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ORGANISM: Artificial Sequence
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                                                                                                                                                                      FEATURE:
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13 TAAGGTGAAGAT 2
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2 GAUAACGUGAAG 13
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Malven, Marianne
                                                   Conservative
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                                                                                                                                 Description of Artificial Sequence: fully synthesized primer
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Pred. No. 6
                                                               Score 10.4; DB 6; Pred. No. 6.2e+04;
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Best Local Similarity
"~+~hes 9; Conserv?
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-310-188-80491
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                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: PCT-US03-03662-97
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SOFTWARE: PatentIn ver
SEQ ID NO 97
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Matches 9; Conser
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TITLE OF INVENTION: BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL VIRAL REGULATORY G
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: 47487
CURRENT APPLICATION NUMBER: US/10/310,188
CURRENT FILING DATE: 2002-12-19
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NUMBER OF SEQ ID NOS: 86841
                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 2003-01-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: PCT/US03/03662
CURRENT FILING DATE: 2003-02-06
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TITLE OF INVENTION: Using Short Interfering Nucleic Acid (siNA)
FILE REFERENCE: 02-1005-A (400/083)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: US 60/363,124
PRIOR FILING DATE: 2002-03-11
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                                                                                                                                            ORGANISM: Artificial Sequence
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15 ATAACGTGAAAA 4
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                                                                                                                                                                                                                                                                                                                FILING DATE: 2002-09-09
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 60/406,784
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Beigelman, Leoni
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Pred. No. 6.2e+04
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Pred. No. 6.2e+04
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US-10-310-188-14089
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                                                                ; ORGANISM: Homo sapiens
US-10-310-188-14089
                                                                                                                      FILE REFERENCE: 47487
CURRENT APPLICATION NUMBER: US/10/310,188
CURRENT FILING DATE: 2002-12-19
NUMBER OF SEQ ID NOS: 86841
SOFTWARE: Patentin version 3.1
SEQ ID NO 14089
LENGTH: 19
  Query Match
Best Local Similarity
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APPLICANT: ROSETTABLE GROUP OF NOVEL VIRAL REGULATORY GENERAL INVENTION: BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL VIRAL REGULATORY GENERAL OF INVENTION: USES THEREOF
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SEQ ID NO 336
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SEQ ID NO 82
LENGTH: 20
TYPE: DNA
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SEQ ID NO 73
LENGTH: 20
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APPLICANT: Dobie, Kenneth W.
APPLICANT: Dobie, Kenneth W.
TITLE OF INVENTION: Modulation of Estrogen-Responsive Finger Protein Expression
FILE REFERENCE: RTS-0473
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TITLE OF INVENTION: MULTIPLEX REAL-TIME QU:
FILE REFERENCE: 21108.0009P1
CURRENT APPLICATION NUMBER: PCT/US02/38806
CURRENT FILING DATE: 2002-12-02
CURRENT FILING DATE: 2002-12-02
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                                                                                                                                                                     ORGANISM: Artificial FEATURE:
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Search completed: May 22, Job time : 374.364 secs

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			UG-2001;	16-A	Patent: WO 0159102-A 109 16-AUG-2001;	JOURNAL
			activity	lease	Nucleozymes with endonuclease activity	TITLE
					Breaker, R. and Emilsson, G.	AUTHORS
					1 (bases 1 to 18)	REFERENCE
		•			artificial sequences.	
					synthetic construct	ORGANISM
					synthetic construct.	SOURCE
					•	KEYWORDS
					AX214296.1 GI:15524373	VERSION
					AX214296	ACCESSION
			59102.	WO01	Sequence 109 from Patent WO0159102.	DEFINITION
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                                                                    Barany, F., Zirvi, M., Gerry, N.P., Favis, R. and Kliman, R. Method of designing addressable array for detection of sequence differences using ligase detection reaction patent: WO 0179548-A 5661 25-OCT-2001; CORNELL RESEARCH FOUNDATION, INC. (US)
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6 c 6 g 6 t
       /organism="synthetic construct"
/db_xref="taxon:32630"
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   OS Uniden
PN 199
PD 24-NOV
PF 15-MAY
PR SHIGER
PC C12N15
PC C12N15/00,
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CC Strand
CC Topolo
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                                                                                                                                                                                                                                                                                                                                                                                                              E30812 20 bp DNA linear PAT 18-JUN-
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Patent: JP 1999318468-A
THE KANAGAWA ACADEMY OF
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Shigeru, N. and Hirofumi, S.
                                                                                                                                                                                                                                                                                                                        unidentified
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Strandedness: Single; Topology: Linear;
                                                              C12N1/21, C12N5/10, C12P21/02, C12Q1/68//(C12N5/10, C12R1:91),
                                                                                 C12N1/19
                                                                                             SHIGERU NOGUCHI, HIROFUMI SUEMORI
C12N15/09, A01K67/027, A61K38/00, A61K48/00, C07K14/47, C07K16/18,
                                                                                                                                                                             Unidentified
JP 1999318468-A/6
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                                A61K37/02, C12N5/00, (C12N5/00, C12R1:91)
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/db_xref="taxon:33630"
/note="Hypothetical Probe Sequence"
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Pred. No. 9.2e+04;
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AR057682
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Grimm,S., Stinchcomb,D.T., McSwiggen,J.,
Draper,K.G.
                                                                                                                          1 (bases 1 to 17)
Grimm,S., Stinchcomb,D.T., McSwiggen,J.,
                                                                                                                                                                                                  Sequence 1977 from patent AR057773
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                                                                                             Intercellular adhesion molecule-1 (ICAM-1) ribozymes Patent: US 5837542-A 1977 17-NOV-1998;
                                                                                                                  Draper, K.G.
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Patent: US 5837542-A 1886 17-NOV-1998;
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Location/Qualifiers
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AX010333
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AR115440
AR115440.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ribozyme treatment of diseases or conditions related intercellular adhesion molecule-1 (ICAM-1) Patent: US 6132967-A 1886 17-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unknown
                                                                                                                                                                                                                 Ribozyme treatment of diseases or conditions related to levels of intercellular adhesion molecule-1 (ICAM-1) Patent: US 6132967-A 1977 17-OCT-2000;
                                                                                                                                                                                                                                                                                                                                       AR115531 17
Sequence 1977 from patent US
AR115531
                                                                                                                                                                                                                                                            1 (bases 1 to 17)
Grimm,S., Stinchcomb,D.T., McSwiggen,J.,
                                                                                                                                                                                                                                                                                               Unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unknown
                                                                                                                                                                                                                                                   Draper, K.G.
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9; Conserv
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5 c 7 g
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5 c 7 g
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Pred. No. 1.5e+05;
3; Mismatches 1
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Pred. No. 1.5e+05;
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                                                                                                                                                                                                                                                                                                                                 Patent: WO 0159103-A 633 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                 artificial sequences.

1 (bases 1 to 17)

Blatt,L., Mcswiggen,J. and Chowrira,B.M.
Method and reagent for the modulation and diagnosis
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Sequence 633 from Patent W00159103.
AX215191
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                                                        Sequence 1 from Patent 105557
                                                                                                                                                                                                                                                                                                                                                                                                                                                        synthetic construct. synthetic construct
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Bacteriophage M13mp18
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/db_xref="taxon:28360"
/note="M13mp18 ssDNA"
3 c 9 g 2 t
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/note="Nucleic Acid"
3 c 3 g
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Pred. No. 2e+05;
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AX293459
                                                  Glover, D.M., Yamamoto, R. and Henderson, D. Mus101 and homologues thereof Patent: WO 014802-A 43 05-JUL-2001; Cyclacel Limited (GB)
                                                                                                                                                                                 18 bp
Sequence 43 from Patent W00148202.
AX189338
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10; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        synthetic construct synthetic construct
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unclassified.

1 (bases 1 to 17)
                                                                                                     artificial sequences.
1 (bases 1 to 18)
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3 c 9 g
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/note="umcal-"
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/organism="synthetic construct"
/db_xref="taxon:32630"
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Pred. No. 3.4e+05;
2; Mismatches 2
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Search completed: May 22, 2003, 19:36:43 Job time: 1049.64 secs
                                                                                                                               Query Match 58.9%; Score 10.6; DB 6; Length 20; Best Local Similarity 58.8%; Pred. No. 4.4e+05; Matches. 10; Conservative 3; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match . 58.9%; Score 10.6; DB 6; Length 18; Best Local Similarity 58.8%; Pred. No. 4.4e+05; Matches 10; Conservative 3; Mismatches 4; Indels
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| | | | :|: ||:|| |
17 AAGGCTTATTGGTGCCA 1
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                                                                                                                                                                                                                                                                                                            Unknown.
Unknown.
Unclassified.
1 (bases 1 to 20)
Wu,R.-Y., You,L.-R. and Soong,T.-S.
Wucleotide sequence of two circular SSDNA associated with banana bunchy top virus and method for detection of banana bunchy top
                                                                                                                                                                                                                                                                               Patent: US 5846705-A 6 08-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  AR063454 20 bp
Sequence 6 from patent US 5846705.
AR063454
AR063454.1 GI:5992762
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="Primer"
5 a 5 c
                                                                                                                                                                                                              Location/Qualifiers
1. .20
/organism="unknown"
a 4 c 7 g
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GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

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Minimum DB
Maximum DB
Result
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
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                                                                                    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 20
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18
Query
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/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1983.DAT:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SIDS2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /SIDS2/
                                                                                                                                                                                              /SIDS2/gcgdata/geneseq/geneseqn-embl/NA2000.DAT:*
/SIDS2/gcgdata/geneseq/geneseqn-embl/NA2001A.DAT:*
/SIDS2/gcgdata/geneseq/geneseqn-embl/NA2001B.DAT:*
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                                           SUMMARIES
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r #1 to ampl	6583	18	20	ŗ	10	44
mer A (Group	9585	16	20		10	43
#7 for	V	24	15	5	10	42
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chromosome	4296	24	20	6	0	40
e oligonucl	564	24	20	6	0	39
re oligonucl	ABI93623	24	20		0	38
in recepto	178	22	20	5	0	37
MTHFR antis	050	22	20	<u>,</u>	0	36
imer for insuli	5979	21	20	6.	0	35 5
R primer used	0316	20	20	9	0	34
primer	0598	20	20	6	0	ω ω
nate primer	6674	18	20	6	0	32
nse oligonu	0330	11	19	σ,	0	31
e le	851	24	18	6	0	30
rlin exon F	524	23	18	5	0	29
r for detec	573	20	18	6	0	28
n NOGO Ambe	266	23	17	6	0	27
n NOGO DNAzy	230	23	17	6	0	26
e flk-1 VEGF	306	18	17	6	0	25
flk-1 VEGF	306	18	17	σ.	0	24
k-1 VEGF	90	18	17	6.	0	23
#4. Uniden	136	22	15	6	0	22
probe 3 us	AAV32883	19	15	5	10.2	21
e oligonucl	9433	24	20	7.	0	20
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e caspase 2 a	7490	24	20	7.	10.4	18
strate for HH	Z6288	21	15	7.	0.	17
e 01:	326	24	20	œ	٥.	16
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rRNA er	4106	24	18	œ	0.	14
sophila mus101	B0868	22	18	œ	0.	13
ture oligonucl	I9350	24	20	0	0	12
man NOGO Hammer	K0063	23	17	62.2	11.2	11
M13mp18 ssDNA ampl	AAZ39793	21	17			10

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ALIGNMENTS

RESULT 1 AAS12348

AAS12348 standard; DNA; 18 BP

AAS12348;

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Deoxyribozyme; cytostatic; endonuclease; RNA cleavage; DNA cleavage; gene therapy; plant; fungus; bacteria; mammal; ribozyme; ss.
                WPI; 2001-536526/59
                                Breaker R,
                                                                          08-FEB-2000; 2000US-0181360
31-MAR-2000; 2000US-0193646
                                                                                                                                                     Synthetic
                                                                                                                                                                                              DNA encoding deoxyribozyme #8
                                                 (RIBO-) RIBOZYME PHARM INC (UYYA ) UNIV YALE.
                                                                                                   08-FEB-2001; 2001WO-US04223
                                                                                                                    16-AUG-2001
                                                                                                                                    WO200159102-A2
                                                                                                                                                                                                               21-NOV-2001
                                Beigelman L,
                                                                                                                                                                                                               (first entry)
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Score

Match

Length

В

Description

DNA encoding deoxy

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100.0 71.1 67.8 65.6 65.6 65.6 63.3 63.3

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AAS12348 ABI95945 ABI93941 AAX94010 AAZ44197 ABI93134 AAT53656

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Rat ICAM hammerhea Rat ICAM hammerhea

Hammerhead

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New nucleic acids with endonuclease activity, such as ribozymes

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RNA. The nucleic acids are used to inhibit gene expression and/or cell proliferation, and can be used to treat a disease or condition. More than one nucleic acid can be independently targeted to the same or different sites in a cell. The nucleic acids may be used to study DNA. The modifications to the nucleic acids optimises their catalytic activity and can maintain or enhance their activity. They exhibit a high degree of specificity for RNA. The present sequence represents the coding
               oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful for defection diseases caused by bacterial infectious agents
                                                                                                                                                                                                                                                                                                                                                                                                                                         14-APR-2000; 2000US-197271P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-APR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Capture oligonucleptide Zip ID#3032 oligo #9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABI95945;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABI95945 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        activity,
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                                                                                                                                                                                                                   Designing capture oligonucleotide probes complementary oligonucleotides hybridize
                                                                                                                                                                                                                                                                                                                                            Barany F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                            (CORR ) CORNELL RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         infection;
                                                                                                                                                                      Example 5; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             igase detection reaction;
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                                                                                                                      present invention describes a method (M1) for designing capture
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18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           K-ras;
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                                                                                                                                                                                                                                                                                                                                          Zirvi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     as; PCR primer; probe; capture probe; mutation detection; action reaction; LDR; p53; BRCA1; BRCA2; infectious diseased in the probability of the probability; cogen; tumour suppressor; human papillomavirus; forensic the probability of the pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 4 A; 4 C; 6
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  Listeria monocytogenes
                                                                                                                                                                   300pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                            FOUND INC
                                                                                                                                                                                                                                                                                                                                            Gerry NP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                            Favis R,
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Haemophilus influenza,
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                                                                                                                                                                                                                                                                                                                                            7.
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of RNA or DN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               preferably
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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RESULT 3
ABI93941
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRA1 gene, p53 gene, human papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the particular sites and identifying if ligation of the oligonuclectide probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sets occurred and correlating (using a computer) identified ligation to presence or absence of the target nucleotide sequences. ABI82074 to ABI97546 represent oligonucleotide sequences used in the exemplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Epstein-Barr virus and polio virus, and parasitic infectious agents selected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such
                                                                                                                                                                                                                                                                                                                                                                                                                     ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection: 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Capture oligonucleptide Zip ID#1028 oligo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-FEB-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  as 21 hydroxylase deficiency, Turner Syndrome and obesity defects.
                                                                                                                                Designing capture oligonucleotide probes complementary oligonucleotides hybridize
                                                                                                                                                                                                                                                                     14-APR-2000; 2000US-197271P
                                                                                                                                                                                                                                                                                                   04-APR-2001; 2001WO-US10958
                                                                                                                                                                                                                                                                                                                                25-OCT-2001
                                                                                                                                                                                                                                                                                                                                                              WO200179548-A2
                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABI93941 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              infectious agents e.g. Cryptococcus neoformans, Candida albicans Aspergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cir
                                                                                                     Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
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                                                                      present invention describes a method (M1) for designing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACGGCTTATCGGTGCG
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                                                                                                                                                                             2002-034366/04
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11; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    K-ras;
                                                                                                    5; Fig 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
                                                                                                                                                                                                         Zirvi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 2 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer; probe; capture probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention.
                                                                                                    300pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                         FOUND INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           71.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6 C;
                                                                                                                                                                                                           Gerry NP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ω
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 12.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                           Favis R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       other;
                                                                                                                              for use on a support with little mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .2e+02
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                                                                                                                                                                                                           Kliman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mutation detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  in the exemplification
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                                                                      capture
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                                                                                                                                              to which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is use

for detecting infectious diseases caused by e.g. Salmonella, Listeria monocytogenes and

Haemophilus influenza, bacterial infectious

agents

useful

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 4
AAX94010/c
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Best Local
AAX91991-X97517 represent PCR primers used to amplify open reading frames and other nucleic acid sequences from the genome of Chlamydia pneumoniae (see AAX91990). C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent otitis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAY34584-
                                                                                                                                                                                                                                                                                                                             04-NOV-1998;
21-NOV-1997;
                                                                                                                                                                                Genome sequence of Chlamydia pneumoniae
                                                                                                                                                                                                                        WPI; 1999-357842/30
                                                                                                                                                                                                                                                                                                                                                                                                                   03-JUN-1999
                                                                                                                                                                                                                                                                                           (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                20-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     W09927105-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chlamydia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       vaccine; neutralising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX94010 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AUGGCCUAUCGGUGCGA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pneumoniae
                                                                                                                                               Disclosure; 1912pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 7 A; 6 C; 4 G; 3 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                             98US-0107078
97FR-0014673
                                                                                                                                                                                                                                                                                                                                                                                98WO-IB01890
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        to amplify an ORF of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       67.8%;
64.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     epitope;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 12.2; DB 24; Pred. No. 1.6e+03;
                                                                                                                                               English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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RESULT 5
AAZ44197/c
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RESULT 6
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                      ABI93134;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inducing the differentiation of cerebral nerve tissue. The protein an the gene are expected to be used for the prevention and the treatment of cerebral nerve diseases. AAZ44193-Z44202 represent primers used in
                                                                  ABI93134 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAY35879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae nucleotides sequences can also be used as immunogenic compositions, especially where the vector directs the expression of a neutralising epitope of C. pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                              Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                             protein
                                                                                                                                                                                                                                                                                                                                                                                                                                       the isolation of the cerebral nerve tissue differentiation associat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes a novel murine protein which is capable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 4; Page 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A new protein tissue cell -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-MAY-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Murine cerebral nerve tissue differentiation DNA primer SA1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (KANA-) ZH KANAGAWA KAGAKU (MEIP ) MEIJI MILK PROD CO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 AATGGCCTAACGATG
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                                                                                                                                                                                      TGGTCTCTCGGTGCG
                                                                                                                                                                                                                                  UGGCCUAUCGGUGCG 17
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10; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                             described
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                                                                                                                                                                                                                                                                                                                                                                            BP; 7 A; 7 C; 4 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       participating in the differentiation of cerebral nerve for the prevention and treatment of cerebral nerve tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tissue differentiation; disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98JP-0152027
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                                                                                                                                                                                                                                                                                                                                                                                                                           in the method of
                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22pp; Japanese.
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                                                                                                                                                                                                                                                                                                     65.6%;
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66.7%;
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CO LTD.
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                                                                                                                                                                                                                                                                              4;
                                                                                                                                                                                                                                                                                                     Score 11.8; DB 21
Pred. No. 2.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 11.8;
Pred. No. 2.
                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                           the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              murine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           treatment; primer;
                                                                                                                                                                                                                                                                              Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                               ers used in associated
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                            Gaps
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15-FEB-2002

(first

entry)

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RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                    (1) have melting temperatures within a narrow range. The method is useful CC for detecting infectious diseases caused by bacterial infectious agents CC e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal CC infectious agents e.g. Cryptococcus neoformans, Candida albicans and CC Aspergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cirus, Epstein Barr virus and polio virus, Entamoeba histolytica and Dracunculus CC selected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus CC medinesis. The method is also useful for detecting genetic diseases such CC as 21 hydroxylase deficiency, Turner Syndrome and obesity defects.

CC Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, CC persone, human papillomavirus types 16 and 18 and liver cancers. The cc method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning CC electron microscope and infrared microscope) the support at the CC particular sites and identifying if ligation of the oligonucleotide probe consense or absence of the target nucleotide sequences. ABI82074 to CC ABI97546 represent oligonucleotide sequences used in the exemplification of the present invention.
                         AAT53656
                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; K-ras; PCR primer; probe; capture probe; mutation detect ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious diffection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; fore environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-APR-2001; 2001WO-US10958
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-034366/04
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                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a method (M1) for designing capture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-APR-2000; 2000US-197271P
AAT53656 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CORR ) CORNELL RES
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                                                                                                                                                                                                                  3 UGGCCUAUCGGUGCG
                                                                                                                                                       TGCCCTATCTGTGCG
                                                                                                                                                                                                                                                                               Similarity
9; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5
                                                                                                                                                                                                                                                                                                                                                                                                          20
                                                                                                                                                                                                                                                                                                                                                                                                    BP;
                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                    1 A;
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                                                                                                                                                                                                                                                                                                         65.6%;
                                                                                                                                                                                                                                                                                                                                                                                                    6 C;
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                                                                                                                                                                                                                                                                                                                                                                                                    9
                                                                                                                                                                                                                                                                                                                                                                                                    G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                      Score 11.8;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Favis R,
                                                                                                                                                                                                                                                                                                            .8e+03
                                                                                                                                                                                                                                                                                                                                             DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kliman
                                                                                                                                                                                                                                                                                                                                      24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        77,
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                                                                                                                                                                                                                                                                            Gaps
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Claim

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Page

203;

407pp;

English

Ribozymes having modified bases and methods for use in inhibiting disease related genes

for

producing

them

The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves ICAM-1 mRNA at the nucleotide base position indicated in the DE line. Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and hairpin ribozyme cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and

McSwiggen Sweedler D Woolf T;

D,

Thompson JD,

Tracz D,

Beigelman L, Usman N,

Wincott

Sullivan SM; incott FE;

Dudycz LW;

Stinchcomb

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WPI; 1995-351090/45

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PRX
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PR
XX
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15-APR-1994;
18-MAY-1994;
                                                                                                                                                                                                     23-SEP-1994;
28-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        intercellular adhesion molecule; rel A; tumour necrosis factor; TNF-alpha; respiratory syncytial virus; RSV; bcr-abl; oncogene; translocation; chronic myelogenous leukaemia; CML; cancer; Philadelphia chromosome; inflammation; autoimmune disease; atherosclerosis; myocardial infarction; stroke; restenosis; transplant rejection; rehumatoid arthritis; psoriasis; myocardial ischaemia; Kawasaki disease; septic shock; HIV; myocardial ischaemia; Kawasaki disease; septic shock;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene
                                                                                                                                                                         03-OCT-1994
07-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-MAR-1997
                                                                                                                                               04-NOV-1994
                                                                                                                                                                                                                                                 08-SEP-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                      30-JAN-1995,
23-FEB-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rattus rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Enzymatic nucleic acid; ribozyme;
                                                                                                                10-NOV-1994;
28-NOV-1994;
                                                                                                                                                              11-OCT-1994
                                                                                                                                                                                                                                   23-SEP-1994
                                                                                                                                                                                                                                                                          19-AUG-1994
                                                                                                                                                                                                                                                                                         17-AUG-1994
                                                                                                                                                                                                                                                                                                                      15-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                                                                       29-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              human immunodeficiency virus; acquired
                                                        (RIBO-)
                                                                                                                                                                                                                                                                                                        L6-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ICAM hammerhead ribozyme target sequence (nt. position 2341).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expression;
                                                        RIBOZYME PHARM INC
b DT, Chowrira B, Diren
Karpeisky A, Kisich K,
JA, Modak A, Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                     94US-0311486.
94US-0311749.
94US-0314397.
                                                                                                                                                                                                                                                                            94US-0292620
94US-0293520
                                                                                                  94US-0345516
94US-0357577
                                                                                                                                                                         94US-0316771
94US-0319492
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94US-0245736
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94US-0218934
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                                                                                      94US-0363233
                                                                                                                                94US-0337608
                                                                                                                                              94US-0334847
                                                                                                                                                             94US-0321993
                                                                                                                                                                                                                                               94US-0303039
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   95US-0380734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           downregulation;
                                                                                                                                                                                                                                                                                                        -0291433
              Direnzo A, Draper KG,
h K, Matulic-Adamic J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          trans cleavage; inhibition; interleukin-5; IL-5; ICAM-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immune deficiency syndrome;
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RESULT 8
AAT53740
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                                            07-OCT-1994;
11-OCT-1994;
04-NOV-1994;
                                                                             28-SEP-1994;
03-OCT-1994;
                                                                                                                                                                            06-JUL-1994;
15-AUG-1994;
                                                                                                   23-SEP-1994;
23-SEP-1994;
                                                                                                                        08-SEP-1994;
                                                                                                                                   02-SEP-
                                                                                                                                                                                                                                  07-APR-1994;
                                                                                                                                                                                                                                                                                                   23-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                         Rat ICAM hammerhead ribozyme target sequence (nt.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 synthesised with modifications that improve their nuclease resistance. The ribozymes cleave the ICAM-1 target sequences and thereby inhibit ICAM-1 expression, making them useful for reducing transplant rejection and alleviating symptoms in patients with rheumatoid arthritis, asthma and other inflammatory disorders.
                                                                                                                                              19-AUG-1994;
                                                                                                                                                        17-AUG-1994
                                                                                                                                                                    16-AUG-1994;
                                                                                                                                                                                                    18-MAY-1994;
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                                                                                                                                                                                                                                                        29-MAR-1994;
                                                                                                                                                                                                                                                                            30-JAN-1995;
                                                                                                                                                                                                                                                                                                                                               W09523225-A2
                                                                                                                                                                                                                                                                                                                                                                   Rattus rattus
                                                                                                                                                                                                                                                                                                                                                                                                                   atherosclerosis; myocardial infarction; stroke; restendernsplant rejection; rheumatoid arthritis; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Philadelphia chromosome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Intercellular adhesion molecule; rel A; tumour necrosis factor; TNF-alpha; respiratory syncytial virus; RSV; bcr-abl; oncogene; translocation; conconic myelogenous leukaemia; CML; cancer; philadalphia bronous.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Enzymatic nucleic acid; ribozyme; trans cleavage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT53740;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT53740 standard; RNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GGCCUAUCGGUGC 16
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                                                                                                                                   1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       l Similarity
12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
94US-0345516.
94US-0357577.
94US-0363233.
                                                                          94US-0314397.
94US-0316771.
                                                                                                                                                                         94US-0271280.
94US-0291932.
                                  94US-0334847.
94US-0337608.
                                                      94US-0321993
                                                                94US-0319492
                                                                                                94US-0311749
                                                                                                                      94US-0303039
                                                                                                                                 94US-0300000
                                                                                                                                            94US-0293520
                                                                                                                                                                                                  94US-0245736
                                                                                                                                                                                                             94US-0228041
                                                                                                                                                                                                                                                       94US-0218934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      downregulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                63.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                         inflammation; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               G; 4 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 11.4; DB 16;
Pred. No. 4.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     interleukin-5;
                                                                                                                                                                                                                                                                                                                                                                                             immune deficiency syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                          septic shock; HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                              restenosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    position 2872).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
       Enzymatic and antisense nucleic acid inhibition of repressor genes useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ribozyme cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease resistance. The ribozymes cleave the ICAM-1 target sequences and thereby inhibit ICAM-1 expression, making them useful for reducing transplant rejection and alleviating symptoms in patients with rheumatoid arthritis, asthma and other inflammatory disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       wrimm S, Karpeisky A, Kisich K, Matulic-Adamic J
McSwiggen JA, Modak A, Pavco P, Beigelman L, Sui
Sweedler D, Thompson JD, Tracz D, Hamer J
                                                                                                                                                12-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents a preferred target sequence an enzymatic nucleic acid (i.e. a ribozyme) which cleaves IC mRNA at the nucleotide base position indicated in the DE lir Regions of the mRNA that do not form secondary folding
                                                                                                                   (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                        11-APR-2000; 2000WO-US09721
                                                                                                                                                                                                    19-OCT-2000
                                                                                                                                                                                                                               WO200061729-A2
                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                  Ribozyme; erythropoietin; granulocyte colony stimulating factor; interferon alpha; ss.
                                                                                                                                                                                                                                                                                                                         Hammerhead ribozyme substrate #299
                                                                                                                                                                                                                                                                                                                                                       16-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                  AAF02004;
                                                                                                                                                                                                                                                                                                                                                                                                           AAF02004 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            structures and that contain potential hammerhead and hairpin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ribozymes having modified bases and methods for use in inhibiting disease related genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1995-351090/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 GGCCUAUCGGUGC 16
                                                               2000-647423/62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGCCUGUCGGUGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
12; Conserv
                                                                                       Zwick M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 1 A; 5 C; 7 G; 4 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                              9908-0129390
                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14
                                                                                         Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             63.3%;
92.3%;
                                                                                                                                                                                                                                                                                                                                                                                                           17
                                                                                                                                                                                                                                                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 11.4;
Pred. No. 4
                                                                                       McSwiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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, Usman N, Wincott FE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .7e+03
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ted in the DE line.
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RESULT 10
AAZ39793
           The invention provides new methods for the analysis of compounds, compounds next cularly nucleic acids, that use a solid support, labeled compounds and mass spectrometry. A novel method of making a set of labeled compounds, using of a support and a set of labels, comprises: (1) at clear one first or intermediate step comprising dividing the support into lots, performing a different chemical reaction on each lot of the support so as either to modify that lot of the support or to couple a chemical. CC support with a different label, and combining the lots of the support. CC support into lots, performing a different chemical reaction on each lot of the support that lot of the support with a different label, that lot of the support or to couple a chemical modely to that lot of the support, tagging a fraction on each lot of the support to that lot of the support, tagging a fraction or each lot of the support with a different label, where each different composed is linked to a chemical modety a labeled compound which is separable from the support, and combining the lots of the support. The controlleds and products are used for the analysis of chemical components.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ39793 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 3 A; 6 C; 5 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-062435/05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid analysis; mass spectrometry; chemical compound analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                    spectrometry
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                                                                                                                                                                                                                                                                                                                                                                                            Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              production of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          duction of erythropoietin, granulocyte colony stimulating and interferon alpha.
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and products
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INNOVATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amplifying primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98EP-0303873
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99WO-GB01561
                                                                                                                                                                                                                                                                                                                                                                                            26; 65pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     164pp;
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62.5%;
  are used
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  for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Housby JN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 w
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              enzymatic and
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  analysis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hamilton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           labeled
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compounds
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The invention relates to a nucleic acid molecule necession of a CD20 gene and a nucleic acid molecule necession of a neurite growth inhibitor gene (NOGO). regulates expression of a neurite growth inhibitor gene (NOGO). The nucleic acids (e.g., a ribozyme or a The nucleic acids (e.g., a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN motif) pr an amberzyme (cleaving RNA with an NGN triplet), a zinzyme motif) pr an amberzyme (cleaving RNA with an NGN triplet), a zinzyme motif). The CD20-targetting nucleic acid is used

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNAzyme;
B-cell ly
                                                                                                                               Nucleic acid molecules, e.g., enzymatic nucleic constructs, which down regulate expression of a growth inhibitor gene useful for treating, e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         particularly nucleic acid
                                                                                                                                                                                                                                  (MCSW/)
                                                                                                                                                                                                                                                                                        11-FEB-2000;
28-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                  chemotherapy-induced neuropathy; amyotrophic lateral sclerosis;
Parkinson's disease; ataxia; Huntington's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        inflammatory arthropathy; central nervous system injury;
cerebrovascular accident; CVA; Alzheimer's disease; mult
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia;
B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia;
human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human NOGO Hammerhead Ribozyme #633
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABK00633;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABK00633 standard;
                                                                                                Claim
                                                                                                                       and
                                                                                                                                                                            WPI;
                                                                                                                                                                                                  Blatt L,
                                                                                                                                                                                                                                                                             06-MAR-2000;
                                                                                                                                                                                                                                                                                                                        09-FEB-2001; 2001WO-US04273.
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                                                                                                                                                                                                                                                      (RIBO-)
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                                                                                                                      central nervous system injury
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AUGGCCUAUCGGUGCG
                                                                                                88; Page 76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       l Similarity
11; Conser
                                                                                                                                                                                                                      ) RIBOZYME PHARM I
) BLATT L.
) MCSWIGGEN J.
) CHOWRIRA B M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17
                                                                                                                                                                                                McSwiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP;
                                                                                                                                                                                                                                                                           ; 2000US-181797P.
; 2000US-185516P.
; 2000US-187128P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA;
                                                                                                200pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    62.2%;
68.8%;
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                                                                                                                                                                                                  Chowrira BM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 11.2;
Pred. No. 6
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                                                                                                                              CD20 gene u
lymphoma,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hammerhead ribozyme
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                                                                                                                                 or neurite, leukemia,
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                                                                                                                                          neurite
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Inwunnodeficiency virus) associated NHL, mantle-cell lymphoma (MCL), immunnodeficiency virus) associated NHL, mantle-cell lymphoma (MCL), immunnocytoma (IMC), small B-cell lymphocytic lymphoma, immunne thrombocytopaenia, and inflammatory arthropathy. The NGGO-targetting nucleic acid is used to cleave RNA of the NGGO gene in the presence of a divalent cation that is preferably Mg^2^+. Furthermore, the nucleic acid may be contacted with a cell to reduce NGGO activity of the cell and treat a patient having a condition associated with the level of NGGO. The Inparticular, the NGGO-targetting nucleic acid may be used to treat in particular, the NGGO-targetting nucleic acid may be used to treat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           to cleave RNA of CD20 in the presence of a divalent cation that is preferably Mg^2^+ Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-Hodgkin's lymphoma B-cell, bulky
Example 5; Fig 29; 300pp; English
                                                    Barany F,
                                                                                                                                                                                                                                                                 (CORR ) CORNELL RES
                                                                                                                                                                                                                                                                                                                         14-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                           04-APR-2001; 2001WO-US10958
                                                                                                                                                                                                                                                                                                                                                                                                                                           25-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200179548-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious diffection, 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; fore
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Capture oligonucleptide Zip ID#588 oligo #9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABI93501;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABI93501 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 7 A; 3 C; 3 G; 4 U; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease, muscular dystrophy, and/or other neurodegenerative states which respond to the modulation of NOGO expression. \ensuremath{\mathsf{T}}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS) parkinson's disease, ataxia, Huntington's disease, Creutzfeldt-Jakob
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               central nervous system (CNS) injury and cerebrovascular accident (CVA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present sequence is a hammerhead ribozyme of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 AAUGGCCUAUCGGUGC 16
                                                                                                                                                     2002-034366/04
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9; Conserv
                                                                                                                                                                                                          Zirvi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                         2000US-197271P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer; probe;
                                                                                                                                                                                                                                                              FOUND INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     62.2%;
56.2%;
                                                                                                                                                                                                       Gerry NP,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 11.2; DB 23; Dred. No. 6.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.
                                                                                                                                                                                                       Favis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                capture probe;
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                                                                                                                                                                                                       Kliman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mutation detection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              forensic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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Qy

The present sequence is a Drosophila mus101 genomic

primer which is used to sequence and partial cDNAs. The mus101 is

the a member of

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RESULT 13
AAD08688/c
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Best Local
                                                                                         Novel mus101 polypeptide, a member of BRCT superfamily derived from Drosophila useful for identifying substance capable of affecting mus101 function, and for treating tumor \, -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer is specifically associated with a gene selected from BRCAl gene, passociated with a gene selected from BRCAl gene, and inference of the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the particular sites and identifying if ligation of the oligonucleotide probe sets occurred and correlating (using a computer) identified ligation to a presence or absence of the target nucleotide sequences. ABI82074 to ABI97546 represent oligonucleotide sequences used in the exemplification of the present invention.
                                                                                                                                                                              WPI; 2001-418282/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Aspergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cirus, Epstein-Barr virus and pollo virus, and parasitic infectious agents selected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects. Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the
                                                    Example; Page 44; 108pp; English
                                                                                                                                                                                                                                                                                                   24-DEC-1999;
                                                                                                                                                                                                                                                                                                                                        21-DEC-2000; 2000WO-GB04956
                                                                                                                                                                                                                                                                                                                                                                                05-JUL-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           e.g. Salmoneila, Listeria monocytogenes and Haemophilus influenza, fi
infectious agents e.g. Cryptococcus neoformans, Candida albicans and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a method (M1) for designing capture oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where
                                                                                                                                                                                                                                                                                                                                                                                                                     WO200148202-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA repair;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus101; BRCA1 C-Terminus; BRCT; gene therapy; tumour; mitosis inhibitor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Drosophila mus101 genomic and partial cDNA sequencing primer, GENX9P8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAD08688 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (I) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents
                                                                                                                                                                                                                                                         (CYCL-) CYCLACEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity nes 10; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 AUGGCCUAUCGGUG 15
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                                                                                                                                                                                                                    DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cell
                                                                                                                                                                                                                    Yamamoto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry
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                                                                                                                                                                                                                                                                                                 99GB-0030708
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cycle regulation; passive immunotherapy; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14
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71.4%;
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70
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                                                                                                                                                                                                                      Henderson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 10.8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20;
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RESULT 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                probe and mus101 polynucleotide in the sample. The mus101 and its polynucleotide are useful in gene therapy. The mus101 is useful for identifying a substance capable of affecting mus101 function, and the substance is useful for treating tumour, for inhibiting mitosis and for increasing the susceptibility of a tumour cell to a DNA damaging agent. The mus101 is also useful for identifying substances which affects DNA repair and cell cycle regulation, in vitro or in vivo cell culture system to study the role of mus101 and its homologues in disease, and as immunogens. The antibody to mus101 is useful in diagnosis and in passive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BRCT (BRCA1 C-Terminus) superfamily. The mus101 polynucleotide probe is used for detecting the presence or absence of mus101 polynucleotide in a biological sample by bringing the biological sample containing DNA or RNA into contact with mus101 polynucleotide probe under hybridising conditions, and detecting any duplex formed between mus101 polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18
                                 The invention relates to specifying complex microbes as an index to evaluate a water treatment process. The method involves: (1) extracting microbial DNA which cannot be adapted; (2) amplifying partial regions of the microbial genes; (3) separating the aggregate of the resultant DNA fragments; and (4) using the DNA to compare the growing conditions of the microbes and conditions of the water treatment process. The present sequence represents a probe specific for a 16s rRNA encoding DNA, used in the course of the invention.
                                                                                                                                                                                                                                                                                                               WPI; 2002-440454/47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JP2002101884-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16s rRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABL41065 standard; DNA; 18
                                                                                                                                                                                                                                                         Specifying microbes for evaluation and process -
                                                                                                                                                                                                                                                                                                                                                                                       27-SEP-2000; 2000JP-0294854
                                                                                                                                                                                                                                                                                                                                                                                                                             27-SEP-2000; 2000JP-0294854
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Microbe; water treatment; 16s rRNA; probe; ss.
                                                                                                                                                                                                               Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immunotherapy. It is also useful for detecting mus101 in a biological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17 AAGGCTTATTGGTGCCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 AUGGCCUAUCGGUGCGA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    encoding DNA specific probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 5 A;
                                                                                                                                                                                                             Page 6; 11pp; Japanese.
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58.8%;
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Pred. No. 1.4e+04;
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                                                                                                                                                                                                                                                                         of waste water
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Query Match Best Local S. Matches 11

Local Similarity hes 11; Conserv

58.98; 5 C; 6

Score 10.6; DB 24; Pred. No. 1.4e+04; 2; Mismatches 4;

Length

18;

Search completed: May 22, Job time: 178.636 secs

2003,

19:04:27

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18

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RESULT 15
AAV71835
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                                           Ş
                                                                                  Matches
                                                                                                  Query Match
Best Local
                                                                                                                                                                                           Sequences AAV71835 to AAV71841 represent PCR primers specific for the banana bunchy top virus (BBTV) DNA. The invention provides BBTV DNA I (clone 7-4-2) circular single stranded (css) DNA, CC and DNA II (clone 2-17) css DNA sequences (AAV71832 and AAV71833) and the proteins (AAW87459 and AAW87460) encoded by the open reading frames in the nucleotide sequences. The nucleic acid sequences (AAV71830 to AAV71833) are used as the basis for the construction of PCR primers, to detect BBTV infection. The PCR technique is used for detecting BBTV in plant tissues (preferably banana, especially Musa species). The virus, one of the most important banana species viruses, causes phloem damage and is transmitted by aphids. PCR detection gives accurate, reliable and specific determination of absence or presence of the virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acids having banana bunchy top virus component sequences used to design primers for use in polymerase chain reaction detection of the virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Banana bunchy top virus; BBTV; PCR technique; plant; tissue; banana Musa; phloem damage; aphid; nucleic acid detection; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bunchy top banana virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAV71835;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 10; Column 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-059037/05.
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GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

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Database :
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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ALIGNMENTS

	JOURNAL COMMENT	TITLE	REFERENCE	ACCESSION VERSION KEYWORDS SOURCE ORGANISM	RESULT 1 AZ514794/c LOCUS DEFINITION
University of Utah Genome Center University of Utah Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT 84112, USA Tel: 801 585 5606 Fax: 801 585 717 Email: ddunn@genetics.utah.edu Insert Length: 10000 Std Error: 0.00	plasmid inserts Unpublished (2000) Contact: Robert B. Weiss	1.Slam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Rellly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Welss, R. and Wright, D., Welss, R. Mouse, whole genome scaffolding with paired end reads from 10kb	<pre>Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 20) Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,</pre>	Clone UUGCIMU36ICI3 K, DNA sequence. AZ514794 AZ514794.1 GI:10696110 GSS. house mouse. Mus musculus	AZ514794 20 bp DNA linear GSS 05-OCT-2000 1M0361C13R Mouse 10kb plasmid UUGC1M library Mus musculus genomic

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                                                                                                                                                                                                                                                                      and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                 Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1M0419H15R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0419H15 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seq primer: CACACAGGAAACAGCTATGACC Class: plasmid ends High quality sequence stop: 20.
                                              Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                   University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus
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                                                                                                                                                                                                                           plasmid inserts
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                                                                                                                                                                                                                                                                                                                                                            Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
                                                                                                                                                                                 Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          nouse mouse.
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                                                                                                                  308,
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                                                                                           USA
                                                                                                             Biomedical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AF129072.1), a copy number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

6 c 7 g 3 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /lab_host="E. Coli strain XL10-Gold, Tl-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Laboratory Mouse DNA Resource
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/clone="UUGC1M0361C13"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /sex="Male"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   organism="Mus musculus"
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                                                                                                         Polymers Research Bldg.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA sequence.
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No. 6.6e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                potato.

Solanum tuberosum

Solanum tuberosum

Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;

Spermatophyta; Magnoliophyta; eudicotyledons; Solanaceae; Solanum.

Asteridae; euasterids I; Solanales; Solanaceae; Solanum.
                                                                                                                                                                                                                                                                                          Unpublished (2002)
Other_ESTs: EST619237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BQ511821 15 1
EST619236 Generation of a set
                                                                 This clone is available through the Research Genetics, contact the Research Genetics for further information 1-800-711-6195 or
                                                                                                                                                                          The Institute for Genomic Research 9712 Medical Center Dr, Rockville,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         analyses mixed potato tissues Solanum tuberosum cDNA 5' end, mRNA sequence.
cdna@resgen.com
Seq primer: T3.
                                                                                                                                                                                                                                                       Contact: Robin Buell
                                                                                                                                                                                                                                                                                                                                                                Generation of a set of potato
                                                                                                                                                                                                                                                                                                                                                                                                                                          Restrepo, S., Griffiths, H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Buell, C.R., Hart, A., Baker, B., Tanksley, S., Fry, W., Smart, C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BQ511821.1
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                                                                                                                                                                                                                                                                                                                                                                                                           Karamycheva,S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Plate: 0419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                           potato@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pMD42 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated that the control of the control
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   with adaptors complementary to the insert adaptors and
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/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Mus musculus"
/strain="C57BL/6J"
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64.3%;
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Pred. No. 1e+06;
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for microarray
A clone STMHU18
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                                                                                                                                                                                                                                                 High
                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0054 row: D column: 06
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                      University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                 Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                              Tel: 801 585 5606 Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dunn, D., Aoyagi, A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                   308, Biomedical
                                                                                                                                                                                                                                               quality sequence stop: 19.
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                                                                                                                                                                                                                                                                                                                                                                                                     USA
                   /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="Vector: pBluescript SK(-); Site_1: EcoRI; Site_2: XhoI; supplier: Combination of untreated and Phytophthora infestans-treated libraries of stolons, leaves, leaflets, axillary buds of stem explants, petioles, germinating eyes
                                                    /lab_host="E. Coli strain xL10-Gold, T1-resistant, F
/note="Yector: PWD42nv; Purified genomic DNA from M.
                                                                                                       /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                  /db_xref="taxon:10090"
/clone="UUGC1M0054D06"
                                                                                                                                                                   /organism="Mus musculus"
/strain="C57BL/6J"
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(http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                               /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eutheria;
                                                                                                                                                                                                                                ocation/Qualifiers
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/clone="STMUTTO"
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5 c 2 g
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Rođentia;
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2.3e+06;
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RESULT 5
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                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0054 row: D column: 06
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dunn,D., Aoyagi,A., Barber,M., Beacorn,T.; Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reil,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern, and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                       High quality sequence stop: 19
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                               Tel: 801 585 5606 Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 19)
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1M0054D06F Mouse 10kb plasmid UUGC1M library
clone UUGC1M0054D06 F, DNA sequence.
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8; Conserv
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University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 house mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AZ515014.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated
/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jack Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Biomedical Polymers Research Bldg.,
                                                                                                                                                   /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                        /sex="Male"
                                                                                                                            /clone-"UUGC1M0054D06"
                                                                                                        /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                           'lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
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3.1e+06;
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                    from the Jackson
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(http://www.jax.org/resources/documents/dnares/). The DNA

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Query Match
Best Local :
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nes 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tissue Procurement: Miklos Palkovits, M.D., Ph.D. cDNA Library Preparation: Michael J. Brownstein (NHGRI), Shiraki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mammalia; Eutheria; Primates;
1 (bases 1 to 20)
VIH-MGC h++- '
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 bp mRNA linear EST 23-JAN-2001
602287066F1 NIH_MGC_95 Homo sapiens cDNA clone IMAGE:4375648 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                         http://image.llnl.gov
Plate: LLAM10041 row: d column:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution: found through the I.M.A.G.E. Consortium/LLNL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            On Jan 16, 2001 this sequence version replaced Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Li constructed by M. Brownstein (NIMH/NHGRI, National
                                                                                            purified.
                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4375648"
/clone_lib="NIH_MGC_95"
                                                                                                                                                                                                                    /tissue_type="hippocampus"
/lab_host="DH10B"
                                                                                                                                                                                                'note="Organ: brain; Vector: pBluescriptR (modified
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pred. No. 3.1e+06;
2; Mismatches 3;
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Note: this is a NIH_MGC Library."
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4 GGCCUAUCGGU 14
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EST631493
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Samac,D.A., Dickman;M., Town,C.D., Van Aken,S., Utto
Cheung,F. and Fraser,C.M.
ESTs from mycelia of Colletotrichum trifolii race l
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 612 625 1243
Fax: 651 649 5058
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Department of Plant Pathology University of Minnesota
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Deborah A. Samac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Other_ESTs: EST631492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: debbys@puccini.crl.umn.edu
TIGR sequence name: MTSAA51TV More information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              495 Borlaug Hall,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Fungi; Ascomycota; Pezizomycotina;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seq primer: (gth AtA CgA CtC ACt AtA ggg C)
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                                                                                                                                                                          /note="Vector: pBluescript SK+; Site_1: EcoRI; Site_2: EcoRI; isolate: 2sp2; cDNA was prepared from polyA+ enriched RNA The cDNA was ligated into Lambda gtl1 from Stratagene and packaged using Gigapack packaging extracts. An aliquot of the amplified library was used to transduce E. coli Y1090 and phage DNA was purified from a liquid lysate. The cDNA inserts were gel purified after EcoRI digestion and ligated into pBluescript SK+. Aliquots of the ligation were used to transform E. coli DH5alpha which
                                                                                                                                                                                                                                                                                                                                            /tissue_type="mycelia"
/dev_stage="Young, actively growing mycelia (3 days after inoculation) grown in liquid culture (cutin minimal mediur containing 2%glucose)."
/lab_host="DH5alpha"
                                                                                                                           recombinants
3 c
                                                                                                                                                           were plated onto medium with X-gal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /strain="race 1"
/db_xref="taxon:5466"
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                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="DSCT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone="pDSCT1-5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               organism="Colletotrichum trifolii"
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RESULT 8
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Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Eukaryota; As Tetrahymenina; Tetrahymena.
                                                                                                                                       Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 19)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
                                                                                                                                                                                                                                                                                                                                                                                        AZ787588 19 bp DNA linear GSS 16 2M0034A07F Mouse 10kb plasmid UUGCIM library Mus musculus clone UUGC2M0034A07 F, DNA sequence.
      University of Utah Genome Center University of Utah Rm. 308, Biomedical Polymers Res
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                                                                  Contact: Robert B.
                                                                                   Unpublished (2000)
                                                                                                                            Mouse whole genome
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Molecular Genetics and Cell Biology
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                                                                                                           inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Tetrahymena thermophila"
/strain="CU448:1"
/db_xref="taxon:5911"
/clone_lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
proc. Natl. Acad. Sci USA, 98: 8709-8713."
a 3 c 8 g 4 t
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54.5%;
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                                                                                                                      scaffolding with paired end reads from 10kb
Polymers Research Bldg.,
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RESULT 10
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                        University of Utah University of Utah
                                                                                         plasmid inserts
Unpublished (2000)
                                                                                                                                 Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Welss,R.
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Insert Length: 10000 Std Error:
Plate: 0034 row: A column: 07
Seq.primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                     Contact: Robert B.
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1 (bases 1 to 19)
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Fax: 801 585 7177
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Biomedical Polymers Research Bldg.,
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (http://www.jax.org/resources/documents/dnares/). The DNA
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/clone="UUGC2M0034A07"
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/strain="C57BL/6J"
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Pred. No. 4.8e+06;
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Insert Length: 10000 Std Error:
Plate: 0058 row: H column: 14
Seq primer: CACACAGGAAACAGCTATGACC
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2M0015I19F Mouse 10kb plasmid UUGCIM library clone UUGC2M0015I19 F, DNA sequence.
  Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutęleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (Dases 1 to 20)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,H., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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Fax: 801 585 7177
                                                                                                   Unpublished (2000)
                                                                                                                       Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
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Biomedical Polymers Research Bldg.,
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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/note="Vector: PWD42nv; Purified genomic DNA from M.
nusculus C57BL/6J (male) was obtained from the Jackson
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/strain="C57BL/6J"
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Pred. No. 4.8e+06;
2; Mismatches 2
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ORGANISM
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Location/Qualifiers
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Plate: 0015 row: I
                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 20)
Chen, H.M., Rossier, C., Chrast, R. and Antonarakis, S.E.
Cloning of trapped exons from human chromosome 21
                                                                                                                                                                                                                                                                                                 Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Laboratory Mouse DNA Resource
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/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GI:1437772
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA for trapped
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Pred. No. 4.9e+06;
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   Morris, M.A., Lalioti, M.D. and
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                                                     ATGGGCACGCGGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymer
Hymenostomatida; Tetrahymenina; Tetrahymena
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antonarakis, S.E.
Cloning of 559 potential exons of genes of human chromosome 21
                                                                                                                                                                                                                                                                                                                                               920 E. 58th Street, Chicago,
Tel: 773 702 4374
Fax: 773 702 3172
                                                                                                                                                                                                                                                                                                                                                                                                                                                 EST
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97011340
                                                                                                                                                                                                                                                                                                                      Seq primer: T3
                                                                                                                                                                                                                                                                                                                                                                                        Molecular Genetics and Cell Biology University of Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Turkewitz
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                                                                                                         Similarity 57.3
8; Conservative
                                                                                                                                                                                                                                                                                                                                Email: apturkew@midway.uchicago.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tetrahymena thermophila.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              from Tetrahymena thermophila,
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                                                                                                                                                                           /clone_lib="Chilcoat/Turkewitz cDNA (large fraction)".
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
3 c 10 g 2 t
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                                                                                                                                                                                                                                                                        organism="Tetrahymena thermophila"
                                                                                                                                                                                                                                                                                                      ocation/Qualifiers
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                                                                                                      Score 7.6;
Pred. No. 6
2; Mismatch
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Pred. No. 4.9e+06;
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     AI625518

AI625518

ty56d06.x1 NCI_CGAP_Ut2 Homo sapiens cDNA clone IMAGE:2283083 3's similar to TR:P93237 P93237 PROLINE-RICH PROTEIN PRP2 PRECURSOR.
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cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 691 Std Error: 0.00
Seq primer: 40m13 fwd. ET from Amersham
High quality sequence stop: 1.
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anal
AI625518
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Tissue Procurement: Christopher Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
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DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
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Contact: Robert Strausberg,
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                      constructed by Bento Soares and M. Fatima Bonaldo 4\ c \ 6\ g \ 4\ t
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/db_xref="taxon:9606"
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Search completed: May 22, 2003, 20:17:07 Job time: 1317.82 secs
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Best Local Similarity
Matches 7; Conserv
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19 GGGTTTTTGGTGCG 6
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Location/Qualifiers
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Unpublished (1997)
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1 (bases 1 to 19)
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CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be

found through the I.M.A.G.E. Consortium/LLNL at:
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Email: ggapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                      /note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: Sall;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.85 kb. Life Technologies catalog #:
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/clone=lib="NCI_CGAP_Uft2"
/tissue_type="moderately-differentiated endometrial
adenocarcinoma, 3 pooled tumors"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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Maximum Match 100%
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  DB
         US-08-584 040-5817

US-09-108-837-3

US-08-948-097-7

US-08-922-177A-108

US-08-872-917-11

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US-08-837-2010-66

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US-08-182-968A-497

US-08-774-306A-497

US-09-064-156A-497

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R APPLICAT R APPLICAT PLICATOR LING DATE: PLICATION LING DATE: PLICATION ME: Warbu ME: Warbu ME: Warbu MC:	STREET: 633 West Fift STREET: Suite 4700 CITY: Los Angeles STATE: California COUNTRY: U.S.A. ZIP: 90071-2066 COUNTRY: DOOR TO STREET COUNTRY: U.S.A. ZIP: 90071-2066 COMPUTER READABLE FORM: MEDIUM TYPE: 3.5" Dis MEDIUM TYPE: Storage COMPUTER: IBM COMPUTER: IBM COMPUTER: IBM SOFTWARE: WORD PERFEC URRENT APPLICATION NUMBER: U APPLICATION UNBER: AUGUST APPLICATION: 435 CLASSIFICATION: 435 BIOD APPLICATION: 435	-620A-1886 e 1886, Application No. 5837542 L INFORMATION: ICANT: Susan Grimm ICANT: Susan Grimm ICANT: Sean T. Stinc ICANT: Sean Sulliva ICANT: Sean Sulliva ICANT: Kenneth G. L E OF INVENTION: REL E OF INVENTION: REL E OF INVENTION: NOISE E OF INVENTION: MOISE E OF INVENTION: MOISE ER OF SEQUENCES: 23 ESPONDENCE ADDRESS: DRESSEE: Lyon & Lyon	28 29 9.6 53.3 20 0 28 9.6 53.3 20 0 0 31 9.6 53.3 20 0 0 33 9.6 53.3 20 0 0 34 9.6 53.3 20 0 0 35 9.6 53.3 20 0 0 37 9.4 52.2 18 0 0 42 9.4 52.2 18 0 0 43 9.4 52.2 18 0 0 45 9.4 52.2 18 0 0 45 9.4 52.2 18 0 0 45 9.4 52.2 18 0 0 45 9.4 52.2 18 0 0 45 9.4 52.2 18 0 0 45 9.4 52.2 18
including application described below: 7008,995 9,1993 7,1992 7,1	ette, 1.44 Mb ette, Dos 5.0 5.1 5.1 708/292,620A , 1994	S/08292620A comb en aper TREATMENT TED TO LEVELS OR CONDITI TED TO LEVELS OR COLLULAR ADHES CULE-1 (I-CAM-1)	2 US-08-488-177-66 2 US-08-481-072A-66 2 US-08-64-336-66 2 US-08-64-336-66 4 US-08-943-731-546 4 US-08-943-731-546 4 US-08-923-589-72 4 US-08-829-581-418-54 4 US-08-829-64A-30 2 US-08-465-380-100 2 US-08-486-397-100 3 US-08-634-641-965-100 4 US-08-326-110A-13 5 US-08-634-641-100 5 US-08-634-641-100

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                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                   CLASSIFICATION DATA:
PRIOR APPLICATION DATA: including
PRIOR APPLICATION DATA: described
PRIOR APPLICATION DATA: 08/08,895
                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                            REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                        FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 633 West Fit
STREET: Suite 4700
CITY: Los Angeles
STATE: California
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                                                                                              (: (213) 955-0440
67-3510
                                          17 base pairs
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Dan T. Stin
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/ENTION: RIBOZYME TREATMENT OF
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                                                                                                                                                                                                                                                                                                                                                                                                                                             IBM Compatible
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linear
                                                                                                                             (213) 489-1600
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92.3%;
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                                                                                                                                                                                                                                                                                                                                                               US/08/292,620A
17, 1994
                                                                                                                                                                                                                                                                                              including application described below:
                                                                              1977:
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                                                                                                               TOPOLOGY:
US-09-071-845-1886
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Best Local Similarity
Matches 12; Conserv
                                                                                                                                                     TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 18
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No.
                                                                     Query Match
Best Local
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                                                       Matches
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TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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COMPOTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 07/9
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STATE: California
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                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
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                           4 GGCCUAUCGGUGC 16
GGCCUGUCGGUGC 14
                                                                     Similarity
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VENTION: RIBOZYME TREATMENT OF
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92.3%;
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                                                                     Pred. No.
                                                                                 Score 11.4;
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                                                    Mismatches
                                                                     7e+02;
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RESULT 4 US-09-071-845-1977

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US-08-418-071-6
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US-09-071-845-1977
            Sequence 6, Application US/08418071 Patent No. 5846705
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Best Local
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SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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Patent No. 6132967
GENERAL INFORMATION:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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ATTORNEY/AGENT INFORMATION:
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ZIP: 900
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67-3510
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633 West Fifth Street
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UMBER: 08/008,895
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                                                                                                                                                                                        Score 11.4;
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; Sequence 497, Application US/08182968A
; Patent No. 5610054
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INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc= "oligonucleotide primer BB-1"
                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ANTI-SENSE:
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ATTORNEY/AGENT INFORMATION:
NAME: Haley Jr., James F
REGISTRATION NUMBER: 27,
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TITLE OF INVENTION: ASSOCIATED WITH BANANA BUNCHY TOP VIRUS AND METHOD FOR DETE
TITLE OF INVENTION: BANANA BUNCHY TOP VIRUS
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OPERATING SYSTEM:
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les 10; Conserv
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1251 Avenue of the Americas
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You, Li-Ru
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SYSTEM: PC-DOS/MS-DOS
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           IBM Compatible
                                                                                                                                                                   Lyon & Lyon
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METHOD AND REAGENT FOR INHIBITING HEPATITIS C
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                                                                                                                                                                                                    VIRUS REPLICATION 497
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Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6; DB 2;
2.1e+03;
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SOFTWARE:

Word Perfect 5.1

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 497
                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                            REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 223/227
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                            PRIOR APPLICATION NUMBER: 08/182,968
APPLICATION NUMBER: 13, 1994
APPLICATION NUMBER: 07/882,888
FILING DATE: May 14, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: 07/88
FILING DATE: 14-MAY-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
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        TELEFAX: (--
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14 ATGGCCTATTGG 3
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                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
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TELEX: 67-3510
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REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 205/277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FILING DATE: 13-JAN
                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                 SOFTWARE:
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                                                                                                                                                                                                                                  FILING DATE:
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                                                                                                            Warburg, Richard J.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Suite 4700
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    633 West Fifth Street
                                                                                                                                                                                                                                                                                 Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Draper, Kenneth G.
VENTION: METHOD AND REAGENT FOR
VENTION: INHIBITING HEPATITIS C
                               (213) 955-0440
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13-JANUARY-1994
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Pred. No. 2
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Query Match
Best Local Similarity
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US-09-064-156A-497/c
                                                                                  US-09-064-156A-497
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                            Matches
                                         Query Match
Best Local :
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                                                                                                                                                                      TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                     FILING DATE: May 14, 199
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J
                                                                                                                                                       SEQUENCE CHARACTERISTICS:
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                          Local Similarity
nes 8; Conserv
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TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/0
FILING DATE: April 21, 1998
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STATE: California
                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                        FILING DATE: December 26, 1996 APPLICATION NUMBER: 08/182,968
                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: December
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TOPOLOGY: li
                                                                                                                          TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                             FILING DATE:
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2 AUGGCCUAUCGG 13
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                                                                                                                                                                                                  (213) 955-0440
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                          Conservative
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METHOD AND REAGENT FO
INHIBITING HEPATITIS
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66.7%;
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                                                                                                                                                                                                                                                                                                    1992
                                         Score 10.4; DB 3; Pred. No. 2.6e+03;
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Pred. No. 2
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                          Mismatches
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                                                        DB 3;
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US-09-059-779-5
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                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                        equence 5815, Application US/08584040 atent No. 6346398
                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                           TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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                                             CORRESPONDENCE ADDRESS:
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APPLICANT: Michael P. Mittmann
                                                                                                                                                                          APPLICANT:
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CITY: Los Altos
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                                                               IUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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E: Lyon & Lyon 633 West Fifth Street Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Ritter, Van Pelt & Yi LLP 4906 El Camino Real, Suite 205
                                                                                                       RESCOBEDO, Jaime
VENTION: METHOD AND REAGENT FOR THE
VENTION: TREATMENT OF DISEASES OR
VENTION: CONDITIONS RELATED TO LEVELS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
                                                                                                                                                                     McSwiggen, James
Stinchcomb, Dan T.
                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                      Pavco, Pamela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SYSTEM: PC-DOS/MS-DOS
Patentin Release #1.0, Version #1.25
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Synthesis of Diverse Probes on a Substrate
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                                                                            GROWTH FACTOR
                                                                                          OF VASCULAR ENDOTHELIAL
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Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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RESULT 11
US-08-584-040-5816
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
7 hassa nairs
              OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5" Di
MEDIUM TYPE: storage
                                                                                                                                                   STREET: 633 West Fit
STREET: Suite 4700
CITY: Los Angeles
STATE: California
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APPLICATION NUMBER:
                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                       FITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                      APPLICANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                     ITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (41-)
TELEFAX: 67-3510
TEX: 67-3510
                                                                                                                                                                                                                                                                                                                                      ITLE OF INVENTION:
                                        COMPUTER: IBM Com
OPERATING SYSTEM:
 APPLICATION NUMBER:
                                                                                                                          COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
les 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
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                                                                                                                                                                                                                        ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Warburg, Richard
REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 'H: 17 base pairs nucleic acid NDEDNESS: single
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                                                                                                                                                                                                      E: Lyon & Lyon
633 West Fifth Street
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Stinchcomb, Dan T.
                                                                                                                                                                                                                                                                                                                                                                                                 Pavco, Pamela
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                                                            IBM Compatible
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                                                                                                                                                                                                                                                                      GROWTH FACTOR
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                                                                                                                                                                                                                                                                                     OF VASCULAR ENDOTHELIAL
                                                                                                                                                                                                                                                                                                                    TREATMENT OF DISEASES OR
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US/08/584,040
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Pred. No. 3
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                                                                                             1.44 Mb
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RESULT 12
US-08-584-040-5817
; Sequence 5817, Application US/08584040
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Best Local Similarity
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                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
          FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
                                                                                                              APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
APPLICATION NUMBER: 60/005,974
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                       CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 UMBER OF SEQUENCES:
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TELEFAX: (213) 955-0440
TELEX: 67-3510
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STATE: California
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                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A. ZIP: 90071-2066
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633 West Fifth Street
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Stinchcomb, Dan T.
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                                                                                                                                                                                                                 Word Perfect 5.1
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N: 514
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489-1600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               8502
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Pred. No. 3.
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5e+03;
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US-08-948-097-7
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; TYPE: DNA
; ORGANISM: Lactobacillus brevis
US-09-108-837-3
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SEQUENCE CHARACTERISTICS:
LENGTH: 17 base models of the property of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Skerra, Arne
APPLICANT: Voss, Selma
TITLE OF INVENTION: Streptavidin Muteins
FILE REFERENCE: HUBR 1119
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 7
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Best Local :
                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/08/948,097C CURRENT FILING DATE: 1997-10-09 EARLIER APPLICATION NUMBER: DE 196 41 876.3 EARLIER FILING DATE: 1996-10-10
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TITLE OF INVENTION: OLIGONUCLEOTIDE FOR DETECTING LACTIC ACID BACTERIA AND
TITLE OF INVENTION: A NEW METHOD FOR JUDGING BY UTILIZING THEM
FILE REFERENCE: SAMII
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                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 17
                                                                                                                                NAME/KEY: CDS
OTHER INFORMATION: Synthesized
                                                                                                                                                                                                                                        ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                               TYPE:
                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                   LENGTH: 12
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TOPOLOGY: linear
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9; Conservative
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55.6%;
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Score 10;
Pred. No.
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Pred. No. 3.5e+03;
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Pred. No. 3.5e+03;
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                            DB 3;
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                            Length 12;
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Best Local Similarity

4.4e+03;

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TYPE: nucleic acid syran: 20 base pairs
TYPE: nucleic acid syraNDEDNESS; double
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
IMMEDIATE SOURCE:
CLONE: mfd19p1
US-08-222-177A-108
Query Mark
Search completed: May 22, 2003, 20:18:25 Job time: 40.7273 secs
                                                                              Db
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US-08-222-177A-108/c
Sequence 108, Application US/08222177A
Patent No. 5582979
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                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                     Matches
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APPLICANT: Weber, James L.

APPLICANT: Weber, James L.

TITLE OF INVENTION: LEMGTH POLYMORPHISMS IN

TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME

NUMBER OF SEQUENCES: 460

CORRESPONDENCE ADDRESS:

ADDRESSEE: DeWitt Ross & Stevens, S.C.

STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21 APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
GOTTHARE: Details PC-DOS/MS-DOS
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| 18 AATGGCCAAAGGGTTAGA 1
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1 ATCGGTGCGA 10
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STATE: Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE:
                                                                                                             1 AAUGGCCUAUCGGUGCGA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (608) 831-2106
                                                                                                                                                   11; Conservative
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61.1%;
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                                                                                                                                                                Score 10; DB 1;
Pred. No. 4.7e+03;
                                                                                                                                               Mismatches
                                                                                                                                               5; Indels
                                                                                                                                                                                Length 20;
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GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Minimum DB seq length: 0 Maximum DB seq length: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OM nucleic - nucleic search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Title:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               828747 seqs, 660231138 residues
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                                                                                                                                                                                                                                                                                                                                                                                                         Published_Applications_NA: *
/cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
                                                                                                                                                                                                                                                                                    //93n2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
//93n2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
//93n2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

c	c	C	a			a			c	c	C	a	O	O	O		0		Result
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53.3	53.3	υ ω	53,3	53.3	54.4	55.6	55.6	56.7	56.7	56.7	57.8			57.8		62.2	62.2	100.0	Query Match
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US-09-961-077-643	US-09-848-754A-3017	US-09-848-754A-1942	US-09-961-077-264	US-09-825-805-712	US-09-825-805-737	US-09-232-785-373	US-09-961-700A-29	US-09-969-373-2070	US-09-780-533A-2662	US-09-780-533A-2300	US-10-068-160-36	US-09-969-373-3464	US-09-274-553D-519	US-09-504-231A-519	US-09-796-081-4	US-09-796-081-3	US-09-780-533A-633	US-09-780-929-98	ID
			•		Sequence 737, App	373	Sequence 29, Appl		2662,	Sequence 2300, Ap	Sequence 36, Appl	Sequence 3464, Ap	Sequence 519, App	Sequence 519, App	Sequence 4, Appli	Sequence 3, Appli	Sequence 633, App	Sequence 98, Appl	Description

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9-74	9-96	3-60	9-60	3-60	3-60	9-84	9-78	9-78	09-2	9-5	0-26	9-92	30-0	9-60	9-60	9-9	9-60	9-87	9-87	US-09-877	US-09-877	US-09-877	9-87	09-7	0-13
7-3	1-70	-99	-991	-990	-99	8-7	30-5	0-5	74-	04 -	57-1	US-09-924-981-5	30-9	-69	- 69	- 69	-69	7-4	US-09-877-478-	7-4	7-4	17-4	17-4	166-	US-10-132-652
US-09-747-377-21	00A-	-801	-801	US-09-866-108-9651	US-09-866-108-9650	US-09-848-754A-620	US-09-780-533A-2661	33A-	US-09-274-553D-135	231A	US-10-267-117-12	31-5	US-10-080-959A-3	373-	373-	373-	US-09-969-373-2727	US-09-877-478-2309	78-1	-478-1	-478-989	-478-988	US-09-877-478-300	US-09-766-399-17	52-1
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ALIGNMENTS

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RESULT 2
US-09-780-533A-633/c
; Sequence 633, Application US/09780533A
; Publication No. US20030060611A1
; GENERAL INFORMATION:
; APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                     밁
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                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence: Enzymatic Nucleic Acid US-09-780-929-98
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US-09-780-929-98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn version 3.0 SEQ ID NO 98 LENGTH: 18
                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 98, Application US/09780929 Patent No: US20020151693A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/780,929
CURRENT FILING DATE: 2001-02-08
PRIOR APPLICATION NUMBER: US 60/181,360
PRIOR FILING DATE: 2000-02-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Breaker, Ronald
APPLICANT: Beigelman, Leo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Beigelman, Leo TITLE OF INVENTION: Nucleic Acid Catalysts with Endonuclease Activity FILE REFERENCE: MBHB00-884-H (500/001)
      APPLICANT:
                      APPLICANT:
                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 126
                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                           Local Similarity 100 nes 18; Conservative
                                                                                                                                                                                                               1 AAUGGCCUAUCGGUGCGA 18
                                                                                                                                                                                                                                                     1 AAUGGCCUAUCGGUGCGA 18
McSwiggen, Jim
Chowrira, Bharat
                                  Blatt, Larry
                                                                                                                                                                                                                                                                                                            100.0%; Score 18; 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                        RESULT 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: synthetic oligonucleotide US-09-796-081-3
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Best Local Similarity
Watches 9; Conserv
                                                                                                                                                                                                                Sequence 4, Application US/09796081 Publication No. US20030045538A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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Publication No. US20030045538A1
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SEQ ID NO 633
                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/780,533A CURRENT FILING DATE: 2001-02-00 PRIOR APPLICATION
                                      CURRENT APPLICATION NUMBER: US/09/796,081
CURRENT FILING DATE: 2001-08-14
                                                                                                                                 APPLICANT: Danaboyina, Ramaiah
APPLICANT: Nadukkudy, Varghese E
APPLICANT: Joshy, Joseph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/796,081
CURRENT FILING DATE: 2001-08-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Nadukkudy, APPLICANT: Joshy, Jose
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/181,797 PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn Ver.
                                                                            FILE REFERENCE:
                                                                                            TITLE OF INVENTION: VIOLOGEN LINKED ACRIDINE BASED MOLECULE AND PROCESS FOR TITLE OF INVENTION: THE PREPARATION THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: VIOLOGEN LINKED ACRIDINE BASED MOLECULE AND PROCESS FOR TITLE OF INVENTION: THE PREPARATION THEREOF FILE REFERENCE: 3108/01342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 6679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Haeberli, Pete
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Danaboyina, Ramaiah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
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nes 9; Conserv
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                                                                            3108/01342
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56.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                62.2%;
56.2%;
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Pred. No. 5
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Pred. No. 5
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Best Local Similarity
Whiches 8; Conserv:
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                                                                                                                                                                                                                                                RESULT 6
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SEQ ID NO 519
LENGTH: 15
TYPE: RNA
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Best Local Similarity 50...
Thes 9; Conservative
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                                                                                                                                                                              GENERAL INFORMATION:
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                           APPLICANT: MACCJAK, Dennis
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF DISEASES OR CONDITIONS
TITLE OF INVENTION: HEPARITIS C VIRUS INFECTION
FILE REFERENCE: rpi 247/282
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 09/257, 608
PRIOR FILING DATE: 1999-02-24
PRIOR APPLICATION NUMBER: 60/100, 842
PRIOR FILING DATE: 1998-09-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: rpi 247/282
CURRENT APPLICATION NUMBER: US/09/504,231A
CURRENT FILING DATE: 2000-02-15
                                                                                                                                                         APPLICANT: Blatt,
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SURRENT APPLICATION NUMBER: US/09/274,553D
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                                                                                                                      McSwiggen, James
Roberts, Beth
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56.2%;
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Pred. No. 1
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PRIOR APPLICATION NUMBER: 09/257,608

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; ORGANISM: Glycine max US-09-969-373-3464
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Best Local Similarity bu...
Thes 8; Conservative
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 1998-09-18
PRIOR APPLICATION NUMBER: 60/083,217
PRIOR FILING DATE: 1998-04-27
NUMBER OF SEQ ID NOS: 3148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 09/760,427
PRIOR FILING DATE: 2001-01-13
PRIOR APPLICATION NUMBER: US 09/855,768
PRIOR FILING DATE: 2001-05-15
CURRENT APPLICATION NUMBER: US/10/068,160
CURRENT FILING DATE: 2002-02-06
PRIOR APPLICATION NUMBER: 60/128,898
PRIOR FILING DATE: 1999-04-12
                                                                          APPLICANT: VERTHELYI, Daniela TITLE OF INVENTION: OLIGODEOXYNUCLEOTIDE AND FILE REFERENCE: 4239-61999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 09/754,853 PRIOR FILING DATE: 2001-01-05
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CURRENT FILING DATE: 2001-10-02
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APPLICANT: Hauge, Brian M.
TITLE OF INVENTION: Soybean SSRs and Methods of Genotyping
FILE REFERENCE: 38-10(52679)A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE:
                                                                                                                                    APPLICANT:
                                                                                                                                              APPLICANT: THE GOVERNMENT OF THE UNITED STATES OF AMERICA, REPRESENTED BY THE APPLICANT: SECRETARY, DEPARTMENT OF HEALTH AND HUMAN SERVICES APPLICANT: KLINMAN, Dennis
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PRIOR APPLICATION NUMBER: 60/100,842
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
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66.7%;
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66.7%;
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Pred. No. 1.7e
3; Mismatches
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Pred. No. 1.7e+04;
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                                                                                          ITS USE TO INDUCE AN IMMUNE RESPONSE
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Best Local Similarity pu...
Shes 8; Conservative
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
SOFTWARE: Patentin version 3.0
SEQ ID NO 2300
NUMBER OF SEQ ID NOS: 6679
SOFTWARE: PatentIn version 3.0
SEQ ID NO 2662
LENGTH: 17
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Sequence 2300, Application US/09780533A
Publication No. US20030060611A1
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LENGTH: 20
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                                                                     APPLICANT: Chowrita, Bharat
APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Method and Reagent for
FILE REFERENCE: MBHB00,878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Chowrita, Bharat
APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHB00,878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
CURRENT FILING DATE: 2001-02-09
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SOFTWARE: PatentIn ver
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ORGANISM: Homo sapiens
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ORGANISM: Artificial Sequence
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66.7%;
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Pred. No. 2
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Pred. No. 1.
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Query Match
Best Local Similarity
Thes 9; Conserv
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SEQ ID NO 2070
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Matches 8; Conserv
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                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 29
                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 29, Application US/09961700A Publication No. US20020187482A1
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Best Local Similarity
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                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/235,029
PRIOR FILING DATE: 2001-09-25
                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/961,700A
CURRENT FILING DATE: 2002-07-23
                                                                                                                                                                                                                                                                                                       APPLICANT: Wahlestedt, Claes
TITLE OF INVENTION: Methods and Means of RNA Analysis
FILE REFERENCE: 13522-003001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 09/754,853
PRIOR FILING DATE: 2001-01-05
PRIOR APPLICATION NUMBER: US 09/760,427
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/969,373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Soybean SSRs and Methods FILE REFERENCE: 38-10(52679)A
                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Liang, Zicai
APPLICANT: Zhang, Hong-Yan
                                                                                     09-961-700A-29
                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 09/855,768
                                                                                                                                                          TYPE: DNA
                                                                                                                                     ORGANISM: Artificial Sequence
                                                                                                     OTHER INFORMATION: Synthetically generated oligonucleotide
                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9;
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53.3%;
                                 50.0%;
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Pred. No. 2
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Pred. No. 2
                                 Score 10; DB 9;
Pred. No. 2.8e+04;
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                   Mismatches
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                                                Length 18;
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; TYPE: DNA
; ORGANISM: Pinus taeda L.
US-09-232-785-373
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US-09-232-785-373/c
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                                   SOFTWARE: Pate
SEQ ID NO 737
SEQ TO NO 737
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Best Local S
Matches 10
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SEQ ID NO 373
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/232,785
CURRENT FILING DATE: 1999-01-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 09/232,884 PRIOR FILING DATE: 1999-01-15
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TITLE OF INVENTION: MICROSATELITE DNA MARKERS AND USES
TITLE OF INVENTION: THEREOF
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                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: MBHB00-831-F (400/009)
CURRENT APPLICATION NUMBER: US/09/825,805
CURRENT FILING DATE: 2001-09-27
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                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 09/301,511
PRIOR FILING DATE: 1999-04-28
PRIOR APPLICATION NUMBER: 09/186,675
PRIOR FILING DATE: 1998-11-04
                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 09/476,387
PRIOR FILING DATE: 1999-12-30
PRIOR APPLICATION NUMBER: 09/474,432
PRIOR FILING DATE: 1999-12-29
                                                                                                                                                                                                                                                                                                                                                          CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: 09/578,223
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                                                                                                               PRIOR FILING DATE: 1997-11-05
                                                                                                                                PRIOR APPLICATION NUMBER: 60/064,866
                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/083,727
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ITTLE OF INVENTION: Nucleotide Triphosphate
ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity nes 10; Conserv
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                                                                         PatentIn version 3.0
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Nelson, C. Dana
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Beaudry, Amber
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Adamic, Jasenka Matulic
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                                                                                           ID NOS: 1558
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Pred. No.
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SOFTWARE: Patentin version
SEQ ID NO 712
LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
US-09-825-805-712
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US-09-825-805-712
; Sequence 712, Application US/09825805
; Sequence 710, No. US20030004122A1
Search completed: May 22, 2003, 21:38:48 Job time: 87.0909 secs
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                                                                                                                                              Query Match 53.3
Best Local Similarity 75.0
Matches 12; Conservative
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Matches 11; Conservative
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APPLICANT: Zinnen, Shawn
ITTLE OF INVENTION: Nucleotide Triphosphate and their Incorporation into Oligonucleot
FILE REFERENCE: MBHB00-831-F (400/009)
FILE REFERENCE: MBHB00-831-F (400/009)
FILE REFERENT APPLICATION NUMBER: US/09/825,805
FILE REFERENT FAPLICATION NUMBER: US/09/825,805
FILE REFERENT FAPLICATION NUMBER: US/09/825,805
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\PPLICANT: Beigelman, Leo\
\PPLICANT: Beaudry, Amber
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5 CCUAGCGGAGCGA 17
                                                                        2 UGCCCCAGCGGUGUGA 17
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APPLICATION NUMBER: 60/064,866
TITING DATE: 1997-11-05
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FILING DATE: 1999-12-29
APPLICATION NUMBER: 09/301,511
FILING DATE: 1999-04-28
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Beaudry, Amber
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Adamic, Jasenka Matulic
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0; Mismatches
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Pred. No. 4
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3.6e+04;
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq
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Perfect score:
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length: 20
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Gapop 10.0 , Gapext 1.0
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1: /cgn2_b/ptodata/1/pna//

2: /cgn2_b/ptodata/1/pna//

3: /cgn2_b/ptodata/1/pna//

4: /cgn2_b/ptodata/1/pna//
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/cgn2_6/ptodata/1/pna/US092_COMB.seq:*
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/cgn2_6/ptodata/1/pna/US094_COMB.seq:*
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/ptodata/1/pna/US100A_COMB.seq:*
/ptodata/1/pna/US100A_COMB.seq:*
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//ptodata/1/pna/US101A_COMB.seq:*
//ptodata/1/pna/US101B_COMB.seq:*
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/ptodata/1/pna/US099A_COMB.seq:
/ptodata/1/pna/US099B_COMB.seq:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /ptodata/1/pna/US088
                                                                                                                                                                                                            /ptodata/1/pna/US097C_COMB.seq:
/ptodata/1/pna/US098A_COMB.seq:
/ptodata/1/pna/US098B_COMB.seq:
                                                                                                                                                                                                                                                                                                                                                                                                                     pptodata/1/pna/US095B_COMB.seq:*
/ptodata/1/pna/US095C_COMB.seq:*
/ptodata/1/pna/US095D_COMB.seq:*
/ptodata/1/pna/US096A_COMB.seq:*
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/ptodata/1/pna/US096C_COMB.seq:
                                                                                                                                                                                                                                                                                    ptodata/1/pna/US097B_COMB.seq
                                                                                                                                                                                                                                                                                                                                                       /ptodata/1/pna/US096D_COMB.seq
                                                                                                                    ptodata/1/pna/US099C_COMB.seq:
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9_COMB.seq:*
0_COMB.seq:*
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65: /cgn2_6/ptodata/1/pna/US6011_COMB.seq:
66: /cgn2_6/ptodata/1/pna/US6012_COMB.seq:
67: /cgn2_6/ptodata/1/pna/US6012_COMB.seq:
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81: /cgn2_6/ptodata/1/pna/US6030_COMB.seq:
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87: /cgn2_6/ptodata/1/pna
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result	Score	Query .Match Length		DB	ID	Description
1	18	100.0	18	30	US-09-780-929-98	Sequence 98, Appl
c 2	12.2	67.8	20	ω	US-09-827-383-650	650,
ω		67.8	20	31	US-09-827-383-803	803,
c 4		67.8	20	31	US-09-827-383A-650	650,
ر ت		67.8	20	ω,	US-09-827-383A-803	
σ	11.8	65.6	16	ب	PCT-US02-25943-8274	8274,
7	11.8	65.6	16	سر	PCT-US02-25943-18265	S
80	11.8	65.6	16	42	US-10-227-565-8274	•
9	11.8	65.6	16	42	US-10-227-565-18265	18265,
c 10	11.8	65.6	20	15	US-09-198-452A-3336	Sequence 3336, A
11	11.4	63.3	20	28	US-09-703-708-17627	
12	11.4	63.3	20	60	US-60-164-320-17627	
13	11.4	63.3	20	62	US-60-183-791-17627	
c 14	11.2		17	21	US-09-546-745A-1992	Sequence 1992, Ap
c 15	11.2		17	30	US-09-780-533A-633	Sequence 633, App
16	11.2	62.2	19	30	US-09-796-081-3	Sequence 3, Appli
c 17	11.2		19	30	US-09-796-081-4	Sequence 4, Appli
c 18	11.2		20	43	US-10-262-666-77	77
19	10.8		15	ب	PCT-US02-25940-16232	623
20	10.8	60.0	15	ب	PCT-US02-25940-19783	
21	700	,	,			

0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0

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RESULT 2
US-09-827-383-650/c
                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence: US-09-780-929-98
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                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: U$/09/780,929
CURRENT FILING DATE: 2001-02-08
PRIOR APPLICATION NUMBER: U$ 60/181,360
PRIOR FILING DATE: 2000-02-08
NUMBER OF SEQ ID NOS: 126
SOFTWARE: Patentin version 3.0
SEQ ID NO 98
LENGTH: 18
                                                                                                                                                                                                                                        Best Loc
Matches
                                                                                                         GENERAL INFORMATION:
                                                                                                                    Sequence 650, Application US/09827383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 98,
                                                                                                                                                                                                                                                                 Query Match
                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Breaker, Ronald
APPLICANT: Bedyelman, Leo
TITLE OF INVENTION: Nucleic Acid Catalysts with Endonuclease Activity
FILE REFERENCE: MBHB00-884-H (500/01)
                                                              APPLICANT: Michael Mittmann
APPLICANT: Macdonald Morri
APPLICANT: Tom Ryder
                        APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Tag Nucleic Acids and Probe Arrays
                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                            TYPE: RNA
                                                                                                                                                                                    Local
REFERENCE: 3108
ENT APPLICATION NUMBER: US/09/827,383
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Similarity 100.0%;
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                                                   David Lockhart
                                                                             Macdonald Morris
                                                                                                                                                                                                                                        Conservative
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US-09-274-5538-519
US-09-274-5538-519
US-09-274-5538-519
US-09-504-2318-519
US-09-611-931-519
US-09-611-931-519
US-09-633-515-400-26
US-09-705-400-61
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US-08-484-735-6
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US-10-227-563-19783
US-10-227-567-1512
US-10-227-567-1775
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US-09-322-909-4
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Pred. No.
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                                                                                                                                                                                                                                                              Length 18;
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Sequence
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Sequence
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Sequence 16232, A
Sequence 19783, A
Sequence 1512, Ap
Sequence 1775, Ap
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Sequence
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497, App
1054, App
519, App
497, App
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11863, A
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                                                                                                                                                                                                                                                                                                                                                                                                       ; NUMBER OF SEQ ID NOS: 2050
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 803
LENGTH: 20
TYPE: DNA
ORGANIZM: Homo sapiens
US-09-827-383-803
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; ORGANISM: Homo sapiens US-09-827-383-650
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Best Local S
Matches 11
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PRIOR APPLICATION NUMBER: US 60/195,585
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 2050
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 650
LENGTH: 20
TYPE: DNA
            PRIOR APPLICATION NUMBER: US 60/195,585
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEO ID NOS: 2050
SOFTWARE: FASTSEO for Windows Version 4.0
SEO ID NO 650
LENGTH: 20
TYPE: DNA
                                                                                                                                                                                                                        Sequence 650, Applic GENERAL INFORMATION:
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Best Local
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                                                                                                                                          APPLICANT: David Lockhart
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Tag Nucleic Acids and Probe Arrays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Tag Nucleic Acids and
FILE REFERENCE: 3108
                                                                                                    FILE REFERENCE: 3108
CURRENT APPLICATION NUMBER: US/09/827,383A
CURRENT FILING DATE: 2001-04-04
                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                              APPLICANT: Michael Mittmann APPLICANT: Macdonald Morri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/195,585 PRIOR FILING DATE: 2000-04-06
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CURRENT FILING DATE: 2001-04-04
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APPLICANT:
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 ORGANISM: Homo sapiens
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David Lockhart
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64.7%;
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64.7%;
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Pred. No. 2e+04;
3; Mismatches
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                                                                                                                                                               ; OTHER INFORMATION: Chromosome = 1
PCT-US02-25943-8274
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                                                                                                                                                                                                                                                                                                          Sequence 874, Application PC/TUS0225943
GENERAL INFORMATION:
APPLICANT: Feldmann, Richard J.; Global Determinants, Inc.
TITLE OF INVENTION: Pseudomonas aeruginosa PA01, complete genome.
FILE REFERENCE: Jim Zegeer Law Offices - 703-684-8333
CURRENT APPLICATION UNMBER: PCT/US02/25943
CURRENT FILING DATE: 2002-08-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 2050
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 803
                                                                                                    Matches
                                                                                                                                Query Match
                                                                                                                                                                                                                                                                  SOFTWARE: Pro
SEQ ID NO 8274
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                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 64158
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Michael Mittmann
                                                                                                                                                                                                                                      LENGTH: 16
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Tag Nucleic Acids and Probe Arrays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
                                                                                                                                                                                                                  ORGANISM: Pseudomonas aeruginosa PA01, complete genome
                                                                                                                                                                                           LOCATION: (802690)..
                                                                                                                                                                                                            FEATURE:
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                                                                                                    Local Similarity
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Pred. No. 3
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Pred. No. 2
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                                                          APPLICANT: Feldmann, Richard J.; Global Determinants, Inc.
TITLE OF INVENTION: Pseudomonas aeruginosa PA01, complete genome.
FILE REFERENCE: Jim Zegeer Law Offices - 703-684-8333
CURRENT APPLICATION NUMBER: US/10/227,565
CURRENT FILING DATE: 2002-08-26
NUMBER OF SEQ ID NOS: 64158
SOFTWARE: Proprietary
SEQ ID NO 18265
                                                                                                                                                                                                                             Sequence 18265, Application US/10227565 GENERAL INFORMATION:
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FILE REFERENCE: Jim Zegeer Law Offices - 703-684-8333
CURRENT APPLICATION NUMBER: US/10/227,565
CURRENT FILING DATE: 2002-08-26
NUMBER OF SEQ ID NOS: 64158
SOFTWARE: Proprietary
SEQ ID-NO 8274
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Best Local
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SOFTWARE: Proprietary
SEQ ID NO 18265
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TITLE OF INVENTION: Pseudomonas acruginosa PA01, complete genome.
FILE REFERENCE: Jim Zegeer Law Offices - 703-684-8333
CURRENT APPLICATION NUMBER: PCT/US02/25943
CURRENT FILING DATE: 2002-08-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Pseudomonas aeruginosa PAO1, FEATURE:
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OTHER INFORMATION: Chromosome =
                    ORGANISM:
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FEATURE:
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              Pseudomonas aeruginosa PA01,
                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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73.3%;
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Pred. No. 3.
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RESULT 12
US-60-164-320-17627
Sequence 17627, Application US/60164320
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US-09-198-452A-3336/c
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SEQ ID NO 17627
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LENGTH: 20
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CURRENT FILING DATE: 2000-11-02
PRIOR ADDITORNOLLING DATE: 2000-11-02
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TITLE OF INVENTION: Chlamydia pneumoniae genomic TITLE OF INVENTION: thereof and uses thereof, in TITLE OF INVENTION: and treatment of infection FILE REFERENCE: 9710-003-999

CURRENT APPLICATION NUMBER: US/09/198,452A

CURRENT FILING DATE: 1998-11-24
                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE:
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OTHER INFORMATION: Chromosome = 1
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Pred. No. 3
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Pred. No. 3
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US-60-183-791-17627
; Sequence 17627, A
                                                      ; OTHER INFORMATION: Description of Artificial Sequence: US-09-546-745A-1992
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LENGTH: 20
TYPE: DNA
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Matches
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SEQ ID NO 1992
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LENGTH: 20
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Best Local
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                                                                                                                                                                                          TITLE OF INVENTION: Regulation of Repressor Genes using Nucleic Acid Molecules FILE REFERENCE: 237/193
CURRENT APPLICATION NUMBER: US/09/546,745A
CURRENT FILING DATE: 2000-04-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Bower, Stanley G.
APPLICANT: Hinkle, Gregory J.
TITLE OF INVENTION: Xanthomonas campestris
FILE REFERENCE: 38-10(15804)A.
CURRENT APPLICATION NUMBER: US/60/164,320
CURRENT FILING DATE: 1999-11-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/
CURRENT FILING DATE: 2000-02-22
                                                                                                                                                                              NUMBER OF SEQ ID NOS: 7043
                                                                                                                                                                                                                                                                                        APPLICANT:
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                                                                                                                                                                                                                                                                                                                   APPLICANT: Ribozyme Pharmaceuticals, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 38-10(15804)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Bower, Stanley G. APPLICANT: Hinkle, Gregory J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 18992
                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 18992
                                                                                       FEATURE
                                                                                                ORGANISM: Artificial Sequence
                                                                                                                    TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Xanthomonas campestris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                  LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TILE OF INVENTION: Xanthemonas campestris Genome
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les 8; Conser
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les 8; Conserv
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                                                                                                                                                               PatentIn version 3.0
                                                                                                                                                                                                                                                        McSwiggen, Jim
                                                                                                                                                                                                                                                                         Pavco, Pam
                                                                                                                                                                                                                                                                                      Zwick, Michael
                                                                                                                                                                                                                                                                                                   Blatt, Larry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                      Score 11.2;
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Pred. No. 66
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Pred. No. 6
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Sequences

Thereof

Indels Length

0;

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20;

Nucleic Acid

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APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
APPLICANT: Chowrira, Bharat
APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHB00, 878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780, 533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
SOPTWARE: Patentin version 3.0
SEQ ID NO 633
LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
US-09-780-533A-633
                                                                                   В
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US-09-780-533A-633/c
; Sequence 633, Application US/09780533A; GENERAL INFORMATION:
Search completed: May 22, 2003, 21:22:05 Job time: 2076.36 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      В
                                                                                                                                                            Query Match
Best Local Similarity 56...
Matches 9; Conservative
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                                                                             1 AAUGGCCUAUCGGUGC 16
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16 AATGATCTATCTGTGC 1
                                                                                                                                                          62.2%; Score 11.2; DB 30; Length 17; 56.2%; Pred. No. 7.8e+04; tive 4; Mismatches 3; Indels
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Result
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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                                                                                                     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Gapop 10.0 , Gapext 1.0
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18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 /cgn2_6/ptodata/2/pna/PCT_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US07_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US07_NEW_COMB.seq:*
                                                                                                                                                                                                                                                  /cgn2_6/ptodata/2/pna/US10_NEW_COMB.seq2:*
/cgn2_6/ptodata/2/pna/US60_NEW_COMB.seq:*
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(Cgn2_6/ptodata/2/pna/US09_NEW_COMB.seq: *
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PCT-US03-04088-255 US-09-827-383B-1775 US-10-289-762-3336 US-10-259-275-56	US-10-367-892-16232 US-10-367-892-19783	US-10-367-892-19783	40 HO 000 110 HOLD	US-10-287-787-23056 US-10-287-787-23057	US-10-287-787-23056 US-10-287-787-23057 PCT-US03-10154-33	US-10-287-787-23056 US-10-287-787-23057 PCT-US03-10154-33 US-10-404-724-33	US-10-287-787-23056 US-10-287-787-23057 US-10-287-787-23057 PCT-US03-10154-33 US-10-404-724-33 US-10-237-016A-39	US-10-287-787-23056 US-10-287-787-23057 US-10-287-787-23057 PCT-US03-10154-33 US-10-404-724-33 US-10-237-0168-39 US-10-237-106-47	US-10-287-787-23056 US-10-287-787-23057 US-10-287-787-23057 PCT-US03-10154-33 US-10-404-724-33 US-10-237-016A-39 US-10-237-016A-39 US-10-237-016-39	US-10-287-787-23056 US-10-287-787-23057 PCT-USO3-10154-33 US-10-404-724-33 US-10-237-016A-39 US-10-237-016-47 US-10-237-016-39 US-10-237-016-39 US-10-287-787-16018 US-10-287-787-72109	US-10-287-787-23056 US-10-287-787-23057 US-10-287-787-23057 PCT-US03-10154-33 US-10-404-724-33 US-10-237-016A-39 US-10-237-016-39 US-10-237-016-39 US-10-237-016-39 US-10-287-787-15018 US-10-287-787-22109 US-09-827-383B-232	US-10-287-787-23056 US-10-287-787-23057 US-10-287-787-23057 PCT-US03-10154-33 US-10-404-724-33 US-10-237-016A-39 US-10-237-016-47 US-10-237-016-9 US-10-237-016-9 US-10-237-016-9 US-10-287-787-16018 US-10-287-787-22109 US-09-827-3838-232 US-10-168-424-43	US-10-287-787-23056 US-10-287-787-23057 PCT-USG3-10154-33 US-10-247-016A-39 US-10-217-106-47 US-10-217-106-47 US-10-217-106-139 US-10-287-787-22109 US-10-287-787-22109 US-10-287-787-232 US-10-287-787-232 US-10-287-787-232 US-10-287-787-232	US-10-287-787-23056 US-10-287-787-23057 PCT-US03-10154-33 US-10-404-724-33 US-10-237-0164-39 US-10-237-016-47 US-10-237-016-39 US-10-237-016-39 US-10-287-787-15018 US-10-287-787-15018 US-10-287-787-22109 US-09-827-383B-332 US-10-168-424-43 US-10-168-424-43 US-09-274-553E-519 US-10-287-787-12676	US-10-287-787-23056 US-10-287-787-23057 PCT-US03-10154-33 US-10-404-724-33 US-10-237-106A-39 US-10-237-106-47 US-10-237-106-9 US-10-237-016-9 US-10-237-016-9 US-10-287-787-16018 US-10-287-787-1232 US-09-827-3838-232 US-09-827-3838-232 US-10-168-424-43 US-10-287-787-12676 US-10-287-787-142
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56.7	56.7	56.7	56.7	56.7	56.7	56.7	56.7	56.7	56.7	56.7	56.7	57.8	57.8		57.8	57.8	57.8	57.8	57.8	57.8
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Sequence 4121		Sequence 19393	Sequence 19392		Sequence 6615,	Sequence 1488,	Sequence 1285,	Sequence 22461	Sequence 14893				Sequence 22, Appl	Sequence 4, Appl	Sequence 73, Appl	Sequence 11547	Sequence 7571,	Sequence 3310,	Sequence 26977	Sequence 20529

ALIGNMENTS

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                                                                                                                                                            ; OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: region PCT-US03-04088-255
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PCT-US03-04088-255
                                                                                                                                                                                                                                                SOFTWARE: PatentIn vers
SEQ ID NO 255
LENGTH: 19
TYPE: RNA
ORGANIEM: Artificial
                                                                  Query Match 65.6
Best Local Similarity 86.7
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT FILING DATE: 2003-04-28
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR FILING DATE: 2002-07-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/406,784
PRIOR FILING DATE: 2002-08-29
PRIOR APPLICATION NUMBER: US 60/408,378
PRIOR FILING DATE: 2002-09-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/363,124
PRIOR FILING DATE: 2002-03-11
PRIOR APPLICATION NUMBER: US 60/386,782
PRIOR FILING DATE: 2002-06-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2002-09-09
PRIOR APPLICATION NUMBER: US 60/440,129
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: PCT/US03/04088 CURRENT FILING DATE: 2003-04-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA) FILE REFERENCE: 02-708-A (400/080)
                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 626
                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                               FEATURE:
3 UGGCCUAUCGGUGCG 17
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                                                                                                                                                                                                                                                                                                                                                                                            2003-01-15
                                                                                          65.6%;
86.7%;
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                                                                  Score 11.8; DB 1; Pred. No. 8.1e+03; Mismatches 2
                                                                                                               DB 1;
                                                                                                             Length 19;
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                                                                                                                                                                                                                                                                                                                                     ; ORGANISM: Chlamydia pneumoniae US-10-289-762-3336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-289-762-3336/c
                                                                                                                        US-10-259-275-56
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SOFTWARE: Microarray Probe Sequence Listing Generator V
SEQ ID NO 1775
                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 3336
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 3336, Application US/10289762 GENERAL INFORMATION:
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Best Local
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                                                                                      Sequence 56, Application US/10259275 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Chlamydia pneumoniae genomic TITLE OF INVENTION: thereof and uses thereof, in TITLE OF INVENTION: and treatment of infection FILE REFERENCE: 9710-003-999

CURRENT APPLICATION NUMBER: US/10/289,762

CURRENT FILING DATE: 2003-03-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/827,383B
CURRENT FILING DATE: 2001-04-04
PRIOR APPLICATION NUMBER: 60/195,585
PRIOR FILING DATE: 2000-04-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                APPLICANT: Lemon, St
APPLICANT: Y1, Mink
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 6849
CURRENT APPLICATION NUMBER: US/10/259,275
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                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
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TTAG Nucleic Acids and Probe Arrays
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                                                     Y1, MinKyung
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                ON: REPLICATION COMPETENT HEPATITIS C VIRUS AND METHODS OF USE 265.0007 0120
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Pred. No. 8.16
3; Mismatches
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Pred. No. 8.1e+03;
2; Mismatches 2;
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US-10-367-892-19783; Sequence 19783; Appl; GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: US 60/171,909
PRIOR FILING DATE: 1999-12-23
PRIOR APPLICATION NUMBER: US 99/747,419
PRIOR APPLICATION NUMBER: US 60/325,236
PRIOR APPLICATION NUMBER: US 60/325,236
PRIOR FILING DATE: 2001-09-27
PRIOR FILING DATE: 2001-11-13
PRIOR FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 73
                                       NUMBER OF SEQ ID NOS: 25502

SOFTWARE: Proprietary
SEQ ID NO 19783

LENGTH: 15
Type: Fair
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LENGTH: 18
TYPE: DNA
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GENERAL INFORMATION:
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Best Local
                                                                                                                          APPLICANT: Feldmann, Richard J.; Global Determinants, In TITLE OF INVENTION: Deinococcus radiodurans R1 complete FILE REFERENCE: Jim Zeger Law Offices - 703-684-8333 CURRENT APPLICATION NUMBER: US/10/367,892 CURRENT FILING DATE: 2003-03-06
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OTHER INFORMATION: Chromosome = 1
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                                     TYPE: DNA
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ORGANISM: Deinococcus radiodurans R1 complete genome, FEATURE:
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Pred. No. 2
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Pred. No. 1
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RESULT 9
US-10-287-787-23057/c
Sequence 23057, Application US/10287787:
GENERAL INFORMATION:
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; OTHER INFORMATION: Chromosome = 1
US-10-367-892-19783
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GENERAL INFORMATION:
APPLICANT: Feldmann, Richard J.; Global Determinants,
TITLE OF INVENTION: Caulobacter crescentus complete granter REFERENCE: Jim Zegeer Law Offices - 703-684-8333
CURRENT APPLICATION NUMBER: US/10/287,787
CURRENT FILING DATE: 2003-03
NUMBER: OF SEQ ID NOS: 27958
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CURRENT FILING DATE: 2002-11-26
NUMBER OF SEQ ID NOS: 17608
SOFTWARE: Patentin version 3.1
SEQ ID NO 1079
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Pred. No. 3
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Pred. No. 2
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US-10-404-724-33/c
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                                                                                 Sequence 33, Application US/10404724
GENERAL INFORMATION:
APPLICANT: HORWITZ, Arnold H.
TITLE OF INVENTION: Methods and Materials For
TITLE OF INVENTION: Polypeptides
FILE REFERENCE: 13698US01
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Matches
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GENERAL INFORMATION:
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               CURRENT APPLICATION NUMBER: US/10/404,724
CURRENT FILING DATE: 2003-03-31
PRIOR APPLICATION NUMBER: US 60/368,530
PRIOR FILING DATE: 2002-03-29
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CURRENT FILING DATE: 2003-03-31
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NAME/KBY: misc_feature
OTHER INFORMATION: ING-1 Heavy Chain Oligos Low Risk Primers
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TYPE: DNA
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OTHER INFORMATION: Chromosome = 1
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                                                                                                                                                                                                                                                          15 AAGGGCCCATCGGT 2
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nes 9; Conser
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SEQ ID NOS: 79
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                                                                                                                                                                                                                                                                                                                            Conservative
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64.3%;
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Pred. No. 3
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Pred. No. 3
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GENERAL INFORMATION:
APPLICANT: OHTSUBO, KENICHI
APPLICANT: NAKAMURA, SUMIKO
APPLICANT: MIYAMURA, TSUYOSHI
APPLICANT: KUMO, SATOSHI
APPLICANT: KATO, IKUNOSHIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: SYNTHETIC DNA US-10-237-016A-39
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                                                                                                                                                                                                          US-10-217-106-47/c
; Sequence 47, Application US/10217106
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SEQ ID NO 39
LENGTH: 18
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                                                TITLE OF INVENTION: Method of detecting the presence or TITLE OF INVENTION: and identifying the mixed varieties
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TITLE OF INVENTION: DNA-LEVEL RICE PALATABILITY EVALUATION METHOD, AND METHOD OF SEL
TITLE OF INVENTION: PALATABLE RICE THROUGH ANALYSIS OF HALF GRAIN OF UNHULLED/UNPOL
FILE REFERENCE: 227527USOX
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OTHER INFORMATION: ING-1 Heavy Chain Oligos Low Risk Primers -Reverse Primer GR
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17 AATCGCATATCGGT 4
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NUMBER: US/10/217,106: 2002-08-13
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57.1%;
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Pred. No. 3
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                                                  SOFTWARE: Proprietary
SEQ ID NO 16018
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; OTHER INFORMATION: SYNTHETIC DNA US-10-237-016-39
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US-10-237-016-39/c
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Best Local Similarity
Matches 8; Conserv
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                                                                                                          GENERAL INFORMATION:
                                                                                                                           Sequence 16018,
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                  TITLE OF INVENTION: Caulobacter crescentus complete genome FILE REFERENCE: Jim Zegeer Law Offices - 703-684-8333 CURRENT APPLICATION NUMBER: US/10/287,787 CURRENT FILING DATE: 2003-03-03
CURRENT FILING DATE: 2003-03-03
NUMBER OF SEQ ID NOS: 27958
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CURRENT FILING DATE: 2002-09-09
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                                                                                         APPLICANT: Feldmann,
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HARAGUCHI, KAZUTO
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                                                                                           Richard J.; Global Determinants,
                                                                                                                                                                                                                                                                                                          60.0%;
57.1%;
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57.1%;
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Pred. No. 3
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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Gapop 10.0 , Gapext 1.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /cgn2_6/ptodata/1/pna/US06_COMB.seq:*
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n2_6/ptodata/1/pna/US095C_COMB.seq;
n2_6/ptodata/1/pna/US095C_COMB.seq;
n2_6/ptodata/1/pna/US095B_COMB.seq;
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n2_6/ptodata/1/pna/US096E_COMB.seq;
n2_6/ptodata/1/pna/US097B_COMB.seq;
n2_6/ptodata/1/pna/US097B_COMB
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2_6/ptodata/1/pna/US090_COMB.seq:*
2_6/ptodata/1/pna/US091_COMB.seq:*
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2_6/ptodata/1/pna/US093_COMB.seq:*
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/ptodata/1/pna/US083_COMB.seg:*
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44: /cgn2_6/ptodata/1/pna/US6000_COMB.seq: 45: /cgn2_6/ptodata/1/pna/US6001_COMB.seq: 445: /cgn2_6/ptodata/1/pna/US6001_COMB.seq: 447: /cgn2_6/ptodata/1/pna/US6003_COMB.seq: 48: /cgn2_6/ptodata/1/pna/US6005_COMB.seq: 49: /cgn2_6/ptodata/1/pna/US6005_COMB.seq: 50: /cgn2_6/ptodata/1/pna/US6005_COMB.seq: 51: /cgn2_6/ptodata/1/pna/US6005_COMB.seq: 52: /cgn2_6/ptodata/1/pna/US6001_COMB.seq: 53: /cgn2_6/ptodata/1/pna/US6010_COMB.seq: 55: /cgn2_6/ptodata/1/pna/US6011_COMB.seq: 55: /cgn2_6/ptodata/1/pna/US6011_COMB.seq: 56: /cgn2_6/ptodata/1/pna/US6011_COMB.seq: 57: /cgn2_6/ptodata/1/pna/US6011_COMB.seq: 58: /cgn2_6/ptodata/1/pna/US6011_COMB.seq: 59: /cgn2_6/ptodata/1/pna/US6011_COMB.seq: 59: /cgn2_6/ptodata/1/pna/US6011_COMB.seq: 60: /cgn2_6/ptodata/1/pna/US6011_COMB.seq: 61: /cgn2_6/ptodata/1/pna/US6011_COMB.seq: 62: /cgn2_6/ptodata/1/pna/US6011_COMB.seq: 63: /cgn2_6/ptodata/1/pna/US6012_COMB.seq: 63: /cgn2_6/ptodata/1/pna/US6012_COMB.seq: 64: /cgn2_6/ptodata/1/pna/US6012_COMB.seq: 66: /cgn2_6/ptodata/1/pna/US6012_COMB.seq: 66: /cgn2_6/ptodata/1/pna/US6021_COMB.seq: 67: /cgn2_6/ptodata/1/pna/US6021_COMB.seq: 67: /cgn2_6/ptodata/1/pna/US6021_COMB.seq: 67: /cgn2_6/ptodata/1/pna/US6022_COMB.seq: 67: /cgn2_6/ptodata/1/pna/US6023_COMB.seq: 67: /cgn2_6/ptodata/1/pna/US60203_COMB.seq: 67: /cgn2_6/ptodata/1/pna/US60203_COMB.seq: 67: /cgn2_6/ptodata/1/pna/US6033_COMB.seq: 67: /cgn2_6/ptodata/1
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Sequence 217, App Sequence 217, App	Sequence 12172, A				Sequence 12603, A	Sequence 50, Appl	Sequence 22, Appl	59		12	4	4		606				Sequence 1704, Ap	Sequence 97, Appl	Description

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US-09-780-929-97
; Sequence 97, Application US/09780929
; GENERAL INFORMATION:
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APPLICANT: Breaker, Ronald
APPLICANT: Beigelman, Leo
TITLE OF INVENTION: Nucleic Acid Catalysts with Endonuclease Activity
FILE REFERENCE: MBHB00-884-H (500/001)
CURRENT APPLICATION NUMBER: US/09/780,929
CURRENT ETLING DATE: 2001-02-08
PRIOR APPLICATION NUMBER: US 60/181,360
PRIOR APPLICATION NUMBER: US 60/181,360
PRIOR FILING DATE: 2000-02-08
NUMBER OF SEQ ID NOS: 126
SOFTWARE: PatentIn version 3.0
SEQ ID NO 97
LENGTH: 15
                                                                              Sequence 1704, Application GENERAL INFORMATION:
                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                           Matches
                                                   APPLICANT: Pavco, Pamela APPLICANT: Roberts, Elisabeth APPLICANT: Jarvis, Thale
              APPLICANT: Coeshott, Claire
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Method and Reagents for the Treatment of Diseases
TITLE OF INVENTION: to Molecules Involved in Angiogenic Reponses
                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE
                                                                                                                                                                                                                                ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                   TYPE: RNA
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         REFERENCE:
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MBHB00-824-A (239/121)
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4 US-60-201-255-24
6 US-09-996-373-2697
PCT-US02-25942-56
PCT-US02-25942-56
PCT-US02-25942-57
1 US-09-274-553-1004
5 US-09-274-553-469
5 US-09-274-533D-469
9 US-09-504-231B-469
9 US-09-611-931-469
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1 US-09-633-515-447
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US-09-546-745A-5653
                                                                                                                                                                                 Score 15; DB 30;
Pred. No. 3.8e+02;
                                                                                                                                                                                                                       of Artificial Sequence:
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                                                                                                                                                                           Mismatches
                                                                                                                                                                                          Length 15
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Sequence 55, Appl
Sequence 57, Appl
Sequence 447, App
Sequence 447, App
Sequence 469, App
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222, App
223, App
562, App
24, Appl
2697, Ap
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                           or Conditions
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APPLICANT: Zagursky, Robert John
APPLICANT: Metcalf, Benjamin J.
APPLICANT: Metcalf, Benjamin J.
APPLICANT: Peek, Joel A.
APPLICANT: Peek, Joel A.
APPLICANT: Peek, Joel A.
APPLICANT: Fletcher, Leah Diane
TITLE OF INVENTION: CHLAMYDIA TRACHOMATIS GEI
TITLE OF INVENTION: AND POLYPEPTIDES, FRAGMEI
TITLE OF INVENTION: INFECTION
FILE REFERENCE: 9710-0004-999
CURRENT FILING DATE: 198-11-30
CURRENT FILING DATE: 1998-11-30
CURRENT FILING DATE: 1998-11-04
PRIOR APPLICATION NUMBER: US 60/107,077
PRIOR APPLICATION NUMBER: FR 97-16034
PRIOR FILING DATE: 1997-11-217
PRIOR APPLICATION NUMBER: FR 97-15041
PRIOR APPLICATION NUMBER: FR 97-15041
PRIOR FILING DATE: 1997-11-28
NUMBER OF SEQ ID NOS: 5981
NUMBER OF SEG ID NOS: 5981
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3093
LENGTH: 20
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                                                    Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 1998-03-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH:
                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                               FEATURE:
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ATAACGTGGAGAT
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Sankaran, Banumathi
Sankaran, Banumathi
Fletcher, Leah Diane
Fletcher, Leah Diane
FLETCHER, FRACHOMATIS GENOMIC SEQUENCE
TYPENTION: CHLAMYDIA TRACHOMATIS GENOMIC SEQUENCE
TYPENTION: CHLAMYDIA TRACHOMATIS FRACMOSIS, PREVENTION AND TREATMENT OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hoiseth, Susan
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                                                                       Conservative
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69.2%;
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Pred. No.
                                                                                      Score 11.4; DB 1
Pred. No. 4.7e+04
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                                                                       Mismatches
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2.1e+04;
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US-10-003-919-50/c
US-10-003-919-50/c
; Sequence 50, Application US/10003919
; GENERAL INFORMATION:
; APPLICANT: C. Frank Bennett

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; OTHER INFORMATION: PCR primer
US-09-579-536C-4
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                                 US-09-688-078-12
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                                                                                                                                                     NUMBER OF SEQ ID NOS: 22
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 12
   Query Match
                                                                                                                                                                                                                                                                                          APPLICANT: Kask, I
APPLICANT: Melch
APPLICANT: Chin,
                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 4, Application US/09579536C GENERAL INFORMATION:
APPLICANT: MACIAG, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local
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                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/688,078
CURRENT FILING DATE: 2000-10-13
PRIOR APPLICATION NUMBER: US 60/159,622
PRIOR FILING DATE: 1999-10-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: PCT/US97/09407
PRIOR FILING DATE: 1997-05-30
PRIOR APPLICATION NUMBER: US 60/018,841
PRIOR FILING DATE: 1996-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/579,536C CURRENT FILING DATE: 2000-05-24 PRIOR APPLICATION NUMBER: US 09/199,865 PRIOR FILING DATE: 1998-11-25
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SOFTWARE: PatentIn versi
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TITLE OF INVENTION: PROTEINS AND NUCLEIC ACIDS
FILE REFERENCE: 053689-5002-01
                                                                                                                                                                                                                                                           NPPLICANT: Melcher, Thorsten
NPPLICANT: Chin, Daniel J.
PITLE OF INVENTION: DIAGNOSTIC MARKER FOR NEUROLOGICAL CONDITIONS
FILE REFERENCE: 019488-001310US
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                                                                                           TYPE: DNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
                                                                      OTHER INFORMATION: Description of Artificial Sequence: PCR primer
                                            OTHER INFORMATION: EF-hand primer
                                                                FEATURE:
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SMALL, Deena
PRUDOVSKY, Ig
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71.4%;
Score 10.8;
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Pred. No. 16
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Pred. No. 1
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Length 18;
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                                        NAME/KEY:
LOCATION:
                                                                                                     LENGTH: 18
TYPE: DNA
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OTHER INFORMATION: Description of Artificial Sequence: cDNA; OTHER INFORMATION: amplification 5'-primer US-10-213-329-4
                                                                                                             SOFTWARE: Pace
SOFTWARE: Pace
SEQ ID NO 5964
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Best Local Similarity
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                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/60/216,745
CURRENT FILING DATE: 2000-06-30
NUMBER OF SEQ ID NOS: 13665
                                                                                                                                                                                                                                 FILE REFERENCE:
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PRIOR APPLICATION NUMBER: PCT/US97/09407
PRIOR FILING DATE: 1997-05-30
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CURRENT EILING DATE: 2002-08-06
PRIOR APPLICATION NUMBER: US/09/199,865
PRIOR FILING DATE: 1998-11-25
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APPLICANT: Maciag, Thomas
OTHER INFORMATION: upstream amplification primer 99-25537 for SEQ 1433
                                                                          ORGANISM: Homo Sapiens
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Abderrahim, Hadi
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Pepper, Michael S.
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71.4%;
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2; Mismatches
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Pred. No. 1
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US-09-730-289B-606/c Sequence 606, Application US/09730289B GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; NAME/KEY: primer_bind ; LCATION: 1..19 CHATION: 1..19 ; OTHER INFORMATION: downstream amplification primer 99-50547 for US-60-216-745-13493
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CURRENT FILING DATE: 2001-12-06
NUMBER OF SEQ ID NOS: 87
SEQ ID NO 50
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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Best Local Similarity 84.6%;
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PRIOR APPLICATION NUMBER: US 60/169,100
PRIOR FILING DATE: 1999-12-06
NUMBER OF SEQ ID NOS: 3897
SOFTWARE: PatentIn version 3.0
SEQ ID NO 606
                                                                                      APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for Treatment of Cardiac Disease
FILE REFERENCE: Method and Reagent for Treatment of Cardiac Disease
CURRENT APPLICATION NUMBER: US/09/730,289B
CURRENT FILING DATE: 2000-12-05
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CURRENT FILING DATE: 2000-06-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo Sapiens
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les 9; Conserv
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Abderrahim, Hadi
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Pred. No. 7.
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7.8e+04;
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US-09-730-289B-607/c
US-09-579-536-4
                                                                   SEQ ID NO 4
LENGTH: 18
TYPE: DNA
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Best Local Similarity
Matches 10; Conserv
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LENGTH: 17
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Best Local S
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                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1998-1 PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                              TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC METHODS AND COMPOSITIONS TITLE OF INVENTION: BASED ON JAGGED/NOTCH PROTEINS AND NUCLEIC ACIDS FILE REFERENCE: 36-1U2 (200036.0010) Maciag et al.
CURRENT APPLICATION NUMBER: US/09/579,536
CURRENT FILING DATE: 2000-05-24
                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
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                                                                                                                                                         PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 1996-05-31
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TITLE OF INVENTION: Method and Reagent for Treatment
FILE REFERENCE: MBHB00-864-A (400/006)
                                                                                                                         SOFTWARE:
                                                                                                                                        NUMBER OF SEQ ID NOS:
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                                                                                                                                                                                                PRIOR FILING DATE:
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                                                  ORGANISM: Artificial Sequence
              OTHER INFORMATION: Description of Artificial Sequence: PCR primer
                                 FEATURE:
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